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11786171 PASCAL No.: 94-0663659
Cloning an **ipt** gene from Agrobacterium tumefaciens:
characterisation of **cytokinins** in derivative transgenic **plant**tissue

MCKENZIE M J; JAMESON P E; POULTER RUSSELL T M Univ. Otago, botany dep., Dunedin, New Zealand Journal: Plant growth regulation, 1994, 14 (3) 217-228

Language: English

12/3,AB/138 (Item 6 from file: 144) DIALOG(R)File 144:Pascal (c) 2002 INIST/CNRS. All rts. reserv.

11737745 PASCAL No.: 94-0605454

Transgenic tobacco plants that overproduce cytokinins show increased tolerance to exogenous auxin and auxin transport inhibitors YI LI; XIANGYANG SHI; STRABALA T J; HAGEN G; GUILFOYLE T J Univ. Missouri, dep. biochemistry, Columbia MO 65211, USA Journal: Plant science: (Limerick), 1994, 100 (1) 9-14 Language: English

Transgenic tobacco plants expressing the Agrobacterium tumefaciens cytokinin biosynthetic ipt gene under the control of an auxin-inducible SAUR (Small Auxin-Up RNA) gene promoter were used to study interactions between exogenously applied auxins or auxin transport inhibitors and endogenously produced cytokinins. The transgenic plants used in this study had cytokinin levels about 10-fold higher than non-transformed tobacco plants. In aseptic culture, the transgenic tobacco plants exhibited increased tolerance to the toxic effects of high concentrations of exogenously applied auxins. This tolerance is exemplified by increased plant height and fresh weight in transgenic plants treated with auxin compared to similarly treated non-transformed plants

12/3,AB/139 (Item 7 from file: 144) DIALOG(R)File 144:Pascal (c) 2002 INIST/CNRS. All rts. reserv.

10505359 PASCAL No.: 93-0014610
Altered morphology in transgenic tobacco plants that overproduce

cytokinins in specific tissues and organs
YI LI; HAGEN G; GUILFOYLE T J

Univ. Missouri, dep. biochemistry, Columbia MO 65211, USA Journal: Developmental biology, 1992, 153 (2) 386-395 Language: English

An auxin-inducible bidirectional promoter from the soybean SAUR gene locus was fused to a reporter gene in one direction and a cytokinin biosynthetic gene in the opposite direction and the expression of these fused genes was examined in transgenic tobacco. The Escherichia coli uidA gene, which encodes the enzymes beta -glucuronidase (GUS), was used as the reporter gene and the Agrobacterium tumefaciens ipt gene, which encodes the enzyme isopentenyl transferase, was used as the cytokinin biosynthetic gene

12/3,AB/140 (Item 8 from file: 144) DIALOG(R)File 144:Pascal (c) 2002 INIST/CNRS. All rts. reserv.

08398240 PASCAL No.: 88-0398994

Expression of an Agrobacterium Ti plasmid gene involved in

cytokinin biosynthesis, is regulated by virulence loci and induced by plant phenolic compounds JOHN M C; AMASINO R M

Univ. Wisonsin-Madison, coll. agricultural life sci., Madison WI 53706-1569, USA

Journal: Journal of Bacteriology, 1988, 170 (2) 790-795 Language: ENGLISH

12/3,AB/141 (Item 1 from file: 357) DIALOG(R)File 357:Derwent Biotech Res (c) 2002 Thomson Derwent & ISI. All rts. reserv.

0232193 DBA Accession No.: 99-02294 PATENT A new construct to express phytohormones in developing fruit transgenic plant construction via Agrobacterium tumefaciens-mediated isopentenyl-transferase and tryptophan-2,3-dioxygenase gene transfer and expression AUTHOR: Li Y

CORPORATE SOURCE: Manhattan, KS, USA.

PATENT ASSIGNEE: Univ.Kansas-State-Res.Found. 1998

PATENT NUMBER: WO 9849888 PATENT DATE: 981112 WPI ACCESSION NO.:

99-034673 (9903)

PRIORITY APPLIC. NO.: US 45725 APPLIC. DATE: 970506 NATIONAL APPLIC. NO.: WO 98US9013 APPLIC. DATE: 980506 LANGUAGE: English

ABSTRACT: construct containing either an isopentenyl-A DNA transferase (734 amino acids) or a tryptophan-2,3-dioxygenase (EC-1.13.11.11) (241 amino acids) operably linked to an ovary or developing fruit-specific plant-expressible promoter (e.g (749 bp), AGL (1,051 bp) or PLE36 promoter) is new. Also claimed: Agrobacterium tumefaciens LBA 4404-transformed transgenic plant e.g. tomato (Lycopersicon esculentum), cucumber (Cucumis sativus), watermelon (Citrullus lanatus), tobacco (Nicotiana tabacum), apple (Malus sp.), citrus, pear (Pyrus domestica), fig (Ficus carica), currant, muskmelon, squash, cherry (Prunus sp.), sweet potato (Ipomoea batatas), grapevine (Vitis vinifera), sugarbeet (Beta vulgaris), tea (Camellia sinensis), strawberry (Fragaria sp.), blackberry (Rubus ulmifolius), blueberry (Vaccinium sp.), raspberry (Rubus idaeus), loganberry, rose (Rosa sp.), chrysanthemum, or aubergine (Solanum melongena); a method for producing the transgenic **plant**; and a transgenic seed/embryo. The construct is used to integrate cytokinin /auxin biosynthesis enzymes, to produce seedless fruit in the absence of pollination. (27pp)

12/3,AB/142 (Item 2 from file: 357) DIALOG(R) File 357: Derwent Biotech Res (c) 2002 Thomson Derwent & ISI. All rts. reserv.

0220044 DBA Accession No.: 98-01641 PATENT Vector for insertion of target gene into plants along with a marker gene - tobacco transgenic plant construction for use in crop improvement

AUTHOR: Sugita K; Uesugi M; Matsunaga E; Ebinuma H CORPORATE SOURCE: Tokyo, Japan.

PATENT ASSIGNEE: Nippon-Paper 1997

PATENT NUMBER: WO 9742334 PATENT DATE: 971113 WPI ACCESSION NO.:

97-558990 (9751)

PRIORITY APPLIC. NO.: JP 9780821 APPLIC. DATE: 970331 NATIONAL APPLIC. NO.: WO 97JP1569 APPLIC. DATE: 970509

LANGUAGE: JA

ABSTRACT: A new bacterium (especially Agrobacterium sp.) or virus (especially gemini virus, etc.) vector for the efficient introduction

of a target gene into plants contains a marker gene, preferably a gene involved in the retention of Agrobacterium tumefaciens such as a cytokinin synthesis gene, especially the isopentenyltransferase (ipt) gene from T plasmid DNA, which can be deleted before or after expression of the target gene by application of an external stress such as light, heat or chemical treatment. The deletion can be detected by a morphological change in the transgenic plant tissue. The method is useful for crop improvement, especially for tobacco (Nicotiana tabacum) In an example, plasmid pNPI206 was constructed using beta-galactosidase (EC-3.2.1.23) from plasmid pBI121 as the target gene and ipt as the deletable marker. The ends of the eliminable section were sequences derived from plasmid pNPI128. Plasmid pNPI206 was inserted into A. tumefaciens LBA4404 and used to transform tobacco leaves. The transformants were regenerated in the presence of acetosyringin. (44pp)

12/3,AB/143 (Item 3 from file: 357)
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0219018 DBA Accession No.: 98-00615 PATENT

New genetic constructs for transformation of organisms, particularly plants - Cre-recombinase or Flp-recombinase co-expression with a ribozyme, antisense RNA, sense suppression RNA or plant growth factor biosynthetic gene, in a tobacco or potato transgenic plant

AUTHOR: Surin B P; de Feyter R C; Graham M W; Waterhouse P M; Keese P K; Shahjahan A

CORPORATE SOURCE: Campbell, Australian Capital Territory, Australia; Acton, Australian Capital Territory, Australia.

PATENT ASSIGNEE: CSIRO; Univ.Australian-Nat. 1997

PATENT NUMBER: WO 9737012 PATENT DATE: 971009 WPI ACCESSION NO.: 97-526087 (9748)

PRIORITY APPLIC. NO.: AU 969031 APPLIC. DATE: 960329 NATIONAL APPLIC. NO.: WO 97AU197 APPLIC. DATE: 970327 LANGUAGE: English

ABSTRACT: A new construct has a DNA cassette with a recombinase unit (with a Cre-recombinase or Flp-recombinase gene, terminator and 1st promoter) linked to a transgene unit (with 1 or more transgenes and 2nd promoters), flanked by 2 recombinase-binding recombination loci (e.g. lox or frt sites). The transgene may encode a ribozyme, antisense RNA, co-suppression RNA, or may be a selectable marker, reporter gene or an auxin or cytokinin biosynthetic gene or regulatory sequence (e.g. an ipt gene). An intron may be inserted to disrupt recombinase expression. The cassette may be expressed in a tobacco (Nicotiana tabacum), potato (Solanum tuberosum), sweet potato, Jerusalem artichoke, taro, yam, eucalyptus, pine, aspen, gerbera, chrysanthemum, orchid, lily, rose, fuchsia, azalea, carnation, camellia, gardenia, orange, lemon, grapefruit, tangerine, lime, apple, pear, strawberry, raspberry, loganberry, blackberry, sugarcane, banana, plantain, pineapple or asparagus transgenic plant. The
construct may be used for selective removal or integration of transgenes, with tight regulation of expression. (84pp)

12/3,AB/144 (Item 4 from file: 357)
DIALOG(R)File 357:Derwent Biotech Res
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0204055 DBA Accession No.: 96-14826 PATENT
Use of senescence-associated gene promoters - gene promoter SAG12 and SAG13 and isopentenyl-transferase gene expression in transgenic plant, for application in delayed senescence, and

increased flower, seed, and fruit induction

AUTHOR: Amasino R M; Gan S

CORPORATE SOURCE: Madison, WI, USA.

PATENT ASSIGNEE: Wisconsin-Alumni-Res. Found. 1996

PATENT NUMBER: WO 9629858 PATENT DATE: 961003 WPI ACCESSION NO.:

96-454877 (9645)

PRIORITY APPLIC. NO.: US 413135 APPLIC. DATE: 950329 NATIONAL APPLIC. NO.: WO 96US2313 APPLIC. DATE: 960220

LANGUAGE: English

ABSTRACT: The following are claimed: 1) a genetic construct comprising a senescence-associated gene (SAG)12 or SAG13 promoter sequence operably connected to a protein-encoding DNA sequence not natively connected to the promoter sequence; 2) a cell containing a construct as in 1); 3) a plant containing a construct as in 1); 4) a genetic construct comprising a SAG12 or SAG13 promoter operably linked to a DNA sequence encoding isopentenyl-transferase (IT, EC-2.5.1.8) which catalyzes synthesis of a cytokinin; 5) a transgenic plant with delayed senescence, comprising in it's genome, 5' to 3', a genetic construction including a senescence-associated promoter and a coding region for IT; and 6) a transgenic plant having delayed senescence characteristics, comprising in it's genome a foreign genetic construction which comprises 5' to 3' a senescence-specific promoter, a protein region for IT, and a transcriptional termination sequence, where the foreign gene construction is expressed in tissues entering senescence to delay the senescence of the plant tissues. Such transgenic plants will vegetatively grow longer, producing more flowers, seeds, or fruit. (38pp)

12/3,AB/145 (Item 5 from file: 357)
DIALOG(R)File 357:Derwent Biotech Res
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0199851 DBA Accession No.: 96-10031

Marker-free transgenic **plants** produced by a novel transformation method 'MAT vector system' - multi-auto-transformation vector system with a tRNA-isopentenyltransferase selectable marker and excision by homologous recombination (conference abstract)

AUTHOR: Sugita K; Yamakado M; Ebinuma H CORPORATE AFFILIATE: Nippon-Paper-Ind.

CORPORATE SOURCE: Central Research Laboratory, Nippon Paper Industries, Co., Ltd., 5-21-1, Oji, Kita-ku, Tokyo 114, Japan.

JOURNAL: Plant Physiol. (111, 2, Suppl., 165) 1996

ISSN: 0032-0889 CODEN: PLPHAY

CONFERENCE PROCEEDINGS: Plant Biology '96; 1996 Annual Meeting of the American Society of Plant Physiologists, San Antonio, TX, 27 July-2 August, 1996.

LANGUAGE: English

ABSTRACT: A new transformation method, multi-auto-transformation (MAT) vector system, was developed. MAT-vectors contained a chimeric 35Sgene (tRNA-isopentenyltransferase (EC-2.5.1.8)cytokinin biosynthesis gene) used as the selectable marker. Transgenic shoots were identified as ESP (extreme shooty phenotype) without apical dominance. A site-specific-recombination system (plasmid pSR1) of Zygosaccharomyces rouxii was used in the MAT-vector system (plasmid pNPI132) to remove the ipt gene. After selection of transgenic plants, removal of 35S-ipt genes was detected by appearance of normal shoots from ESPs. In an evaluation experiment with tobacco (Nicotiana tabacum), 48 ESP lines were selected and cultured. Normal elongated shoots appeared in 10 ESPs after 2 mth, and another 19 lines after 4 mth. Shoots from these 29 lines (60%) were normally elongated and rooted. These individuals were confirmed as marker-free transgenic plants by DNA analysis. The 35S-ipt gene was

used as a selectable marker to obtain marker free transgenic

plants in hybrid aspen (Populus sieboldii x Populus
grandidentata). (0 ref)

12/3,AB/146 (Item 6 from file: 357)
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0174253 DBA Accession No.: 95-01074 PATENT

Gene construct for conferring enhanced insect resistance to plants Agrobacterium tumefaciens isopentenyltransferase gene expression
in a transgenic plant using a potato protease-inhibitor-IIK

promoter

AUTHOR: Smigocki A G; Neal Jr J W

PATENT ASSIGNEE: USDA 1994

PATENT NUMBER: WO 9424848 PATENT DATE: 941110 WPI ACCESSION NO.:

94-357754 (9444)

PRIORITY APPLIC. NO.: US 54985 APPLIC. DATE: 930430 NATIONAL APPLIC. NO.: WO 94US4773 APPLIC. DATE: 940428

LANGUAGE: English

ABSTRACT: A new gene construct confers enhanced insect resistance on plants , and comprises a wound-inducible promoter (from a potato (Solanum tuberosum) protease-inhibitor-II or proteaseinhibitor-IIK gene) fused to an isopentenyltransferase gene from Agrobacterium tumefaciens. The construct may be inserted in plasmid pBI221, plasmid pCaMVNEO, plasmid pUC19, plasmid pCMC1100 or plasmid pDG208, or a plant binary vector, e.g. plasmid pEND4K, plasmid pMON120, plasmid pMON200, plasmid pGA472, plasmid pKYLX4, plasmid pKYLX5, plasmid pBIN6, plasmid pBIN19, plasmid pAGS112, plasmid pAGS113 or plasmid pKYLX71 (preferred). The construct may be used to confer insect resistance on a wide variety of plants, including crop plants, fruit trees and ornamentals. In an example, a chimeric cytokinin biosynthetic gene was constructed by fusing a bacterium ipt gene from plasmid pTiB683 to the 5'-regulatory region of the potato PI-IIK gene. An EcoRI-HindIII plasmid pPI-II-ipt fragment was subcloned into plasmid pKYLX71 and mobilized into A. tumefaciens EHA101 (plasmid pEHA101) for infection of Nicotiana plumbaginifolia leaf disks. (25pp)

12/3,AB/147 (Item 7 from file: 357)
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0167299 DBA Accession No.: 94-09850
Transgenic peach plants containing a cytokinin biosynthesis
gene display altered growth in vitro and under greenhouse conditions peach transgenic plant construction via isopentenyltransferase ipt gene expression for increased growth
and compact growth habit (conference abstract)

AUTHOR: Hammerschlag F A; Smigocki A C

CORPORATE AFFILIATE: U.S.Dept.Agr.

CORPORATE SOURCE: USDA/ARS, PSI, Plant Molecular Biology Laboratory,

Beltsville, MD 20705, USA.

JOURNAL: Hortscience (29, 5, 454) 1994

CODEN: HJHSAR LANGUAGE: English

ABSTRACT: Peach (Prunus persica) transgenic plants transformed with the ipt gene from Agrobacterium tumefaciens strain tms328::transposon Tn5 and containing elevated levels of cytokinins were screened in vitro for compact growth habit on 4 different levels of benzyladenine (BA). After 9 wk in vitro, the average number of axillary shoot per plant for 2 of the transformants, 99-1 and 40-1, ranged from 1.5- to 6.6-fold that for the

controls on 0-30 uM BA, whereas the average fresh weight ranged from 1.1- to 3.6-fold that for the controls. 1 Of the transformants, 94-1, produced a greater number of axillary shoots only on 30 uM BA. Rooted through propagation from the original derived transformants were monitored for 30 mth in the greenhouse. The average height of transformants 94-1 and 99-1 after 6 mth in the greenhouse was 88 and 77% of controls, respectively, and after 30 mth was 90 and 75% of control, respectively. In comparison to controls, transformants exhibited a greater number of branches per m per plant after 6 wk, but a reduced number after 30 mth. The introduction of a cytokinin gene may be a useful approach to obtaining peach trees with a compact growth habit. (0 ref)

(Item 8 from file: 357) 12/3,AB/148 DIALOG(R) File 357: Derwent Biotech Res (c) 2002 Thomson Derwent & ISI. All rts. reserv.

0149502 DBA Accession No.: 93-07554 PATENT DNA construct and tuber transgenic plant - cytokinin biosynthesis ipt gene cloning and expression in potato using a plasmid pCGP275 vector for crop improvement

PATENT ASSIGNEE: Calgene 1993 PATENT NUMBER: WO 9307272 PATENT DATE: 930415 WPI ACCESSION NO.:

93-134461 (9316)

PRIORITY APPLIC. NO.: AU 918730 APPLIC. DATE: 911003 NATIONAL APPLIC. NO.: WO 92AU528 APPLIC. DATE: 921002

LANGUAGE: English

ABSTRACT: A new DNA construct contains a plant promoter (e.g. chs) and a sequence (e.g. ipt) encoding a molecule capable of enhancing tuber plant cytokinin levels. The tuber is preferably potato (Solanum tuberosum, preferred), sugarbeet (Beta vulgaris), sweet potato (Ipomoea batatas), onion (Allium cepa), garlic (Allium sativum), artichoke (Cynara scolymus) or Dahlia sp. The construct may be contained in a prokaryote and/or eukaryote vector, capable of integration into the genome or autonomous replication, preferably plasmid pCGP275. The ipt gene may be developmentally regulated, and under the control of an enhancer. The DNA may be used to produce a transgenic plant with 1 or more of the following properties: increased endogenous cytokinin, tuber number and/or wt., stem diameter, height or leaf size; delayed leaf senescence; or increased leaf photosynthetic capacity, leading to increased tuber load The transgenic plant is produced by plasmid yield. Agrobacterium sp., transformation, biolistic mobilization in microprojectile bombardment, microinjection or electroporation. (36pp)

(Item 9 from file: 357) 12/3,AB/149 DIALOG(R) File 357: Derwent Biotech Res (c) 2002 Thomson Derwent & ISI. All rts. reserv.

0116303 DBA Accession No.: 91-03945 Modulating endogenous cytokinin levels - DNA cassette construction for tomato fruit tissue-specific gene expression e.g. during ripening; isopentenyl-transferase gene cloning

and expression in transgenic plant; DNA sequence

PATENT ASSIGNEE: Calgene 1991

PATENT NUMBER: EP 409628 PATENT DATE: 910123 WPI ACCESSION NO.: 91-024190

(9104)

PRIORITY APPLIC. NO.: US 382802 APPLIC. DATE: 890719 NATIONAL APPLIC. NO.: EP 90307925 APPLIC. DATE: 900719

LANGUAGE: English

ABSTRACT: An new expression DNA cassette contains (in 5' to 3' developmentally regulated transcription): a of

transcriptional and translational initiation region; a DNA sequence encoding an enzyme in a cytokinin metabolic pathway, under the transcriptional control of the initiator region; and a transcriptional terminator. At least 1 of the control regions is heterologous to the cytokinin gene. A plant cell containing the DNA cassette, and a method for modification of a plant phenotype using the DNA cassette, are also new. The plant cells are preferably tomato (Lycopersicon esculentum) fruit cells. The cytokinin metabolic pathway is preferably a biosynthetic pathway, and the gene preferably encodes DMA-transferase (isopentenyl-transferase). The developmentally regulated initiation region is preferably from a fruit-specific promoter or an ovary tissue promoter, e.g. the 2All, or Z70 gene. Using the DNA cassette, fruit development, properties, maturation and ripening may be controlled. Other fruits (berries, drupes, hesperidium, pepos) or legume edible portions may also be modified using the DNA cassette. (38pp)

12/3, AB/150 (Item 10 from file: 357)
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0109794 DBA Accession No.: 90-12485

Altering plant morphogenesis by plant genetic engineering transgenic plant construction; tissue-specific gene

expression; hairy root culture and propagation (conference paper)

AUTHOR: Scmuelling T; Schell J; Spena A

CORPORATE AFFILIATE: Max-Planck-Inst.Genet.

CORPORATE SOURCE: Max-Planck-Institut fuer Zuechtungsforschung, D-5000 Koeln 30, Germany.

JOURNAL: BIOTEC-90 (131-36) 1990

CODEN: 9999Y

LANGUAGE: English

ABSTRACT: In vivo genetic manipulation makes it possible to characterize the pleiotropic effects of gene products interacting with normal differentiation mechanisms throughout the life-cycle of a plant, without exogenous plant growth factor application or disrupting the integrity of the plant. Genes which alter plant growth and differentiation may be introduced into the plant gnome and their effects characterized. Exchange of regulatory regions allows altered tissue-specific gene expression. Agrobacterium rhizogenes hairy root culture may be grown in vitro on plant growth factor-free culture medium, and plants may be propagated. Rol gene expression (rolA, rolB and rolC) in plants from hairy root cultures has been studied in detail, and altered morphogenetic The ipt gene of described. characteristics been have tumefaciens encodes an isopentenyltransferase which Agrobacterium causes cytokinin overproduction and developmental alterations in transgenic plants . Better knowledge of regulatory sequences should allow a more accurate targeting of gene expression to specific tissues or development stages. (10 ref)

12/3,AB/151 (Item 1 from file: 434)
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Journal: PLANT CELL, **1989**, V1, N4, P403-413 Language: ENGLISH Document Type: ARTICLE

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08305896 BIOSIS NO.: 000043060894 CYTOKININ BIOSYNTHESIS IN DEVELOPING ZEA-MAYS KERNELS AUTHOR: REINECKE D M; BRENNER M L; RUBENSTEIN I AUTHOR ADDRESS: DEP. PLANT BIOL., UNIV. MINNESOTA, ST. PAUL, MINN. 55108. JOURNAL: ANNUAL MEETING OF THE AMERICAN SOCIETY OF PLANT PHYSIOLOGISTS, PITTSBURGH, PENNSYLVANIA, USA, AUGUST 1-5, 1992. PLANT PHYSIOL (BETHESDA) 99 (1 SUPPL.). 1992. 66. **1992**

CODEN: PLPHA

DOCUMENT TYPE: Meeting RECORD TYPE: Citation LANGUAGE: ENGLISH

1992

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CYTOKININ ANTAGONIST ACTIVITY OF SUBSTITUTED PHENETHYLAMINES IN PLANT CELL CULTURE

AUTHOR: CHRISTOU P; BARTON K A

AUTHOR ADDRESS: AGRACETUS, 8420 UNIVERSITY GREEN, MIDDLETON, WI 53562.

JOURNAL: PLANT PHYSIOL (BETHESDA) 89 (2). 1989. 564-568. 1989

FULL JOURNAL NAME: Plant Physiology (Bethesda)

CODEN: PLPHA

RECORD TYPE: Abstract LANGUAGE: ENGLISH

ABSTRACT: A series of structurally related substituted phenethylamines shows extreme toxicity toward wild-type callus tissue cultures of tobacco (Nicotiana tabacum), soybean (Glycine max), corn (Zea mays), and sunflower (Helianthus annuus L.), but tobacco crown gall cultures are resistant to the compounds. The essential components that result in toxicity of the phenethylamines include one aromatic hydroxyl and one primary aliphatic amino group. A series of attenuated crown gall cultures, generated by transformation of tobacco with various modified Agrobacterium strains, has been used to demonstrate that the resistance of crown galls to the phenethylamines is due to the expression in these tissues of isopentenyl transferase, a first step in cytokinin biosynthesis. The toxicity of the compounds to tissues cultures is very rapid, but can be overcome by prior exposure of the calli to exogenous cytokinin. Because of the relationships we have observed between cytokinins and these compounds, we propose that the substituted phenethylamines may represent a class of chemicals that can be used as specific probes to further an understanding of cytokinin metabolism in plant tissues.

1989

6/3, AB/3(Item 3 from file: 5) DIALOG(R)File 5:Biosis Previews(R) (c) 2002 BIOSIS. All rts. reserv.

04695286 BIOSIS NO.: 000079108415 THE BIOGENESIS OF CYTOKININS

AUTHOR: KLAEMBT D; HOLTZ J; HELBACH M; MAASS H

AUTHOR ADDRESS: BOTANISCHES INST. UNIV., MECKENHEIMER ALLEE 170, D-5300

JOURNAL: BER DTSCH BOT GES 97 (1-2). 1984. 57-66. 1984

FULL JOURNAL NAME: Berichte der Deutschen Botanischen Gesellschaft

CODEN: BEDBA

RECORD TYPE: Abstract

LANGUAGE: GERMAN

ABSTRACT: Cytokinins, N6-substituted adenines, their ribosides and ribotides, act on cell division and cell growth, and are known to delay senescence in leaf explants by attracting amino acids, sugars, phosphate etc. Therefore cytokinins should be involved in growth- and sink-regulation on fruit and storage organs. Since it is known that special tRNA of all different organisms contain these modified nucleotides the assumption arises that cytokinins may be products of tRNA digestion. tRNA half life in Lactobacillus acidophilus and Agrobacterium tumefaciens is 1.5 times the generation time. tRNA in primary roots of Zea mays and Helianthus annuus and in roots of 5 wk old Phaseolus vulgaris posses half lifes of 65 to 75 h. .DELTA.2-Isopentenyldiphosphate: tRNA-.DELTA.2-isopentenyltransferase was prepared from L. acidophilus and Z. mays root tips, caryopses, young and adult leaves. Beside tRNA, MS2-RNA, endogenous oligonucleotides, poly A and oligo A could act as substrates for the isopentenylation. The distribution of the .DELTA.2-isopentenyltranferase in corn showed the highest content related to mg protein in root tips, only 1/10 of that in growing fruits and young leaves and much less in adult leaves. Following up the radiolabeled cytokinins in L. acidophilus after pulse labeling with [14C]-mevalonic acid the [14CC]-cytokinins appeared about 3 h later than the label was incorporated into the tRNA. This is consistent with the hypothesis that tRNA are the source for the free cytokinins. In P. vulgaris fed [14C] -adenine to the roots by pulse labeling and followed up the [14C]-cytokinins in root and leaves as well as the [14C]-incorporation into tRNA and an oligonucleotide fraction and their [14C] -cytokinin-nucleotide content in roots. The profile of [14C]-zeatin in roots and leaves gives no hint for any direct isopentenylation of one of the A-pool derivative but is in complete agreement with the hypothesis describing the cytokinin production by RNA digestion. tRNA account for about 50% only. The residual sources are expected in mRNA, their poly A sequences and/or their degradation products in the form of A containing oligonucleotides.

1984

6/3,AB/4 (Item 1 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
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06611933 Genuine Article#: ZE419 Number of References: 34
Title: Phenotypic alterations and component analysis of seed yield in transgenic Brassica napus plants expressing the tzs gene (
ABSTRACT AVAILABLE)

Author(s): Roeckel P (REPRINT); Oancia T; Drevet JR

Corporate Source: UNIV CLERMONT FERRAND, INRA, LAB ORG & VARIABIL GENOMES VEGETAUX, 24 AVE LANDAIS/F-63177 CLERMONT FERRAND//FRANCE/ (REPRINT); UNIV CALGARY, DEPT BIOL SCI/CALGARY/AB T2N 1N4/CANADA/; UNIV CLERMONT FERRAND, BIOL CELLULAIRE LAB, CNRS, URA 6547 GEEM/F-63177 CLERMONT FERRAND//FRANCE/

Journal: PHYSIOLOGIA PLANTARUM, 1998, V102, N2 (FEB), P243-249

ISSN: 0031-9317 Publication date: 19980200

Publisher: MUNKSGAARD INT PUBL LTD, 35 NORRE SOGADE, PO BOX 2148, DK-1016 COPENHAGEN, DENMARK

Language: English Document Type: ARTICLE

Abstract: Cytokinins play an important role in plant development. We investigated the possibility that the nopaline Ti plasmid gene (tzs) from Agrobacterium tumefaciens could encode a protein able to participate in plant cytokinin production

and lead to alterations in plant phenotype as a result of the expression of endogenous tzs. tzs was placed under the control of a heat-inducible promoter from the Zea mays hsp 70 gene. The expression of this fused gene was examined in transgenic Brassica napus plants. The tzs gene, which encodes the enzyme dimethylallyl transferase, was used as a cytokinin biosynthetic gene. The expression of the tzs gene was monitored by RNA hybridization and analysis of cytokinin. content. Overproduction of cytokinin was observed even when the plants had not been heat-shocked, and the plants displayed a reduced root system, increased height and branching, and delayed flowering. In addition, a significant increase in seed yield was observed in the transgenic plants, accounted for by increased number of seeds per silique and seed weight. The results suggest that increased levels of cytokinins, through the expression of tzs, are correlated with growth rather than with differentiation processes.

6/3,AB/5 (Item 2 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
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06399001 Genuine Article#: YP814 Number of References: 77 Title: Role and function of cytokinin oxidase in plants (ABSTRACT AVAILABLE)

Author(s): Jones RJ (REPRINT) ; Schreiber BMN

Corporate Source: UNIV MINNESOTA, DEPT AGRON & PLANT GENET, 411 BORLAUG HALL, 1991 BUFORD CIRCLE/ST PAUL//MN/55108 (REPRINT)

Journal: PLANT GROWTH REGULATION, 1997, V23, N1-2 (OCT), P123-134

ISSN: 0167-6903 Publication date: 19971000

Publisher: KLUWER ACADEMIC PUBL, SPUIBOULEVARD 50, PO BOX 17, 3300 AA DORDRECHT, NETHERLANDS

Language: English Document Type: ARTICLE

Abstract: Cytokinin oxidase (CK oxidase) is widely distributed in plants and is the only enzyme that has been shown unequivocally to catalyze the catabolism of specific cytokinins (CKs) to inactive products that lack the N-6-unsaturated side chain. Thus, the enzyme is thought to play a major role in controlling the level or species of CKs in plant tissues. However, despite its discovery more than 25 years ago, little attention has been given to the elucidation of its role and function in plant growth and development. This review seeks to bring in to context the current state of knowledge regarding the biochemical and molecular properties, regulation in undifferentiated and differentiated tissues, and recent results from studies using transgenic plants in an attempt to provide a more comprehensive understanding of the physiological significance of the enzyme in plants. Notwithstanding species, tissue and other specific differences, in general, CK oxidase appears to contribute to CK homeostasis in plants. However, complete clarity as to its function awaits purification of the protein to homogeneity and the ultimate development of requisite molecular probes.

6/3,AB/6 (Item 3 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
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06398996 Genuine Article#: YP814 Number of References: 116
Title: Cytokinin conjugation: recent advances and patterns in
plant evolution (ABSTRACT AVAILABLE)

Author(s): Auer CA (REPRINT)

Corporate Source: UNIV CONNECTICUT, DEPT PLANT SCI/STORRS//CT/06269 (REPRINT)

Journal: PLANT GROWTH REGULATION, 1997, V23, N1-2 (OCT), P17-32

number per ear at maturity by up to 30% and in some cases the total kernel weight per ear. The increase was due to a reduction in apical kernel abortion.

6/3,AB/8 (Item 5 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
(c) 2002 Inst for Sci Info. All rts. reserv.

04047799 Genuine Article#: QK335 Number of References: 29
Title: INCREASE OF ENDOGENOUS ZEATIN RIBOSIDE BY INTRODUCTION OF THE
IPT GENE IN WILD-TYPE AND THE LATERAL SUPPRESSOR MUTANT OF TOMATO
(Abstract Available)

Author(s): GROOT SPC; BOUWER R; BUSSCHER M; LINDHOUT P; DONS HJ
COrporate Source: CTR PLANT BREEDING & REPROD RES,CPRO,DLO,DEPT DEV
BIOL,POB 16/6700 AA WAGENINGEN/NETHERLANDS/; CTR PLANT BREEDING &
REPROD RES,CPRO,DLO,DEPT VEGETABLE & FRUIT CROPS/6700 AA
WAGENINGEN/NETHERLANDS/

Journal: PLANT GROWTH REGULATION, 1995, V16, N1 (JAN), P27-36

ISSN: 0167-6903

Language: ENGLISH Document Type: ARTICLE

Abstract: We studied axillary meristem formation of the lateral suppressor (ls) mutant of tomato after elevating the endogenous cytokinin levels through introduction of the isopentenyltransferase (ipt) gene from Agrobacterium tumefaciens. Growth and development of several transformants were examined during in vitro culture. Transformants exhibited phenotypes varying in severity and were divided into four classes. A number of the ipt transformants had a normal phenotype, as non-transformed plants. Others showed a mild to severe 'cytokinin-like' phenotype. Transformants with a mild phenotype exhibited reduced internode length and reduced root development. Transformants with a severe phenotype showed even shorter internodes, loss of apical dominance, reduction of leaf size, production of callus at the basis of the shoots and absence of root development or development of green non-branching roots. The severity of the phenotype correlated well with the level of ipt gene expression, as measured by northern analysis. Transformants with a severe phenotype also exhibited increased levels of zeatin riboside, but zeatin levels were not elevated. The increase in endogenous zeatin riboside levels in the ls mutant did not restore axillary meristem formation, but sometimes bulbous structures were formed in the initially 'empty' leaf axils. Several adventitious meristems and shoots developed from below the surface of these structures. It is concluded that a reduced level of cytokinins in the ls mutant shoots is not responsible for the absence of axillary meristem formation.

6/3,AB/9 (Item 6 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
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03927116 Genuine Article#: QT551 Number of References: 40
Title: THE EFFECT OF AUXIN ON CYTOKININ LEVELS AND METABOLISM IN
TRANSGENIC TOBACCO TISSUE EXPRESSING AN IPT GENE (Abstract Available)

Author(s): ZHANG R; ZHANG X; WANG J; LETHAM DS; MCKINNEY SA; HIGGINS TJV
COrporate Source: AUSTRALIAN NATL UNIV, COOPERAT RES CTR PLANT SCI, POB
475/CANBERRA/ACT 2601/AUSTRALIA/; AUSTRALIAN NATL UNIV, COOPERAT RES CTR
PLANT SCI/CANBERRA/ACT 2601/AUSTRALIA/; AUSTRALIAN NATL UNIV, RES SCH
BIOL SCI, PLANT CELL BIOL GRP/CANBERRA/ACT 2601/AUSTRALIA/; CSIRO, DIV
PLANT IND/CANBERRA/ACT 2601/AUSTRALIA/

Journal: PLANTA, 1995, V196, N1 (MAR), P84-94

ISSN: 0032-0935

Lanquage: ENGLISH Document Type: ARTICLE

ISSN: 0167-6903 Publication date: 19971000

Publisher: KLUWER ACADEMIC PUBL, SPUIBOULEVARD 50, PO BOX 17, 3300 AA DORDRECHT, NETHERLANDS

Language: English Document Type: REVIEW

Abstract: Cytokinin (CK) conjugates are important in plant development because they regulate active CK concentrations, CK transport, storage, and irreversible inactivation. While numerous CK conjugates have been identified in higher plants, the biological functions of these compounds, their location within cells and tissues, and the enzymes and genes involved in their regulation are not clearly understood. In this paper, recent advances are reported which have occurred through the study of transgenic plants containing the ipt or rolC genes, the identification of new regulatory enzymes affecting CKs, and the characterization of new CK conjugates. In addition, a survey of the literature is presented which examines the pattern of CK conjugates found in different plant taxa. Based on current knowledge, it appears that green algae, mosses, and ferns contain relatively few CK conjugates of isopentenyl adenine (iP) and zeatin (Z). In contrast, higher land plants, such as gymnosperms and angiosperms, contain a more complex set of CKs, primarily conjugates of Z and dihydrozeatin (DHZ). This suggests that the pattern of CK conjugation has become more complex in parallel with the increasing complexity of higher plants.

6/3,AB/7 (Item 4 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
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04063789 Genuine Article#: RB577 Number of References: 52
Title: CHANGES IN CYTOKININS AND CYTOKININ OXIDASE ACTIVITY IN
DEVELOPING MAIZE KERNELS AND THE EFFECTS OF EXOGENOUS CYTOKININ
ON KERNEL DEVELOPMENT (Abstract Available)

Author(s): DIETRICH JT; KAMINEK M; BLEVINS DG; REINBOTT TM; MORRIS RO COrporate Source: UNIV MISSOURI, DEPT BIOCHEM, 117 SCHWEITZEL

HALL/COLUMBIA//MO/65211; UNIV MISSOURI, DEPT BIOCHEM/COLUMBIA//MO/65211; ACAD SCI CZECH REPUBL, INST EXPTL BOT/CR-16630 PRAGUE 6//CZECH REPUBLIC/; UNIV MISSOURI, DEPT AGRON/COLUMBIA//MO/65211

Journal: PLANT PHYSIOLOGY AND BIOCHEMISTRY, 1995, V33, N3 (MAY-JUN), P327-336

ISSN: 0981-9428

Language: ENGLISH Document Type: ARTICLE

Abstract: Temporal changes in cytokinin levels, mitotic activity and cytokinin oxidase activity were determined within kernels at the same stage of physiological development in single ears of field-grown maize (Zea mays L.). Cytokinins were qualitatively and quantitatively characterized by immunoaffinity chromatography, high-performance liquid chromatography (HPLC) and radioimmunoassay (RIA). Zeatin (Z), zeatin riboside (ZR) and isopentenyladenosine (iPA) all reached their maximum concentrations 9 days after pollination (DAP). The mitotic activity within the endosperm also peaked at 9 DAP. Cytokinin oxidase was present in kernels at basal levels from 3-6 DAP, then increased substantially through 10 DAP. Comparison of oxidase activity in kernels which are maturing normally and those which will abort, revealed major differences. In aborting apical kernels, the enzyme activity remained at basal levels from 4-10 DAP and only increased slightly trough 15 DAP. In median kernels, which develop normally, oxidase activity increased significantly by 5 DAP and reached a peak 4-fold higher than the basal level by 9 DAP. The differences in cytokinin oxidase activity between kernels which are maturing normally and those which will abort was so pronounced that cytokinin oxidase levels can be considered an indicator of normal kernel development. Stem infusion of benzylaminopurine (BA), but not Z or ZR, into intact plants at pollination increased the kernel

and lead to alterations in plant phenotype as a result of the expression of endogenous tzs. tzs was placed under the control of a heat-inducible promoter from the Zea mays hsp 70 gene. The expression of this fused gene was examined in transgenic Brassica napus plants. The tzs gene, which encodes the enzyme dimethylallyl transferase, was used as a cytokinin biosynthetic gene. The expression of the tzs gene was monitored by RNA hybridization and analysis of cytokinin. content. Overproduction of cytokinin was observed even when the plants had not been heat-shocked, and the plants displayed a reduced root system, increased height and branching, and delayed flowering. In addition, a significant increase in seed yield was observed in the transgenic plants, accounted for by increased number of seeds per silique and seed weight. The results suggest that increased levels of cytokinins, through the expression of tzs, are correlated with growth rather than with differentiation processes.

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Journal: PLANT GROWTH REGULATION, 1997, V23, N1-2 (OCT), P123-134

ISSN: 0167-6903 Publication date: 19971000

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6/3,AB/6 (Item 3 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
(c) 2002 Inst for Sci Info. All rts. reserv.

06398996 Genuine Article#: YP814 Number of References: 116 Title: Cytokinin conjugation: recent advances and patterns in plant evolution (ABSTRACT AVAILABLE)

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Journal: PLANT GROWTH REGULATION, 1997, V23, N1-2 (OCT), P17-32

ISSN: 0167-6903 Publication date: 19971000

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Language: English Document Type: REVIEW

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Author(s): DIETRICH JT; KAMINEK M; BLEVINS DG; REINBOTT TM; MORRIS RO Corporate Source: UNIV MISSOURI, DEPT BIOCHEM, 117 SCHWEITZEL

HALL/COLUMBIA//MO/65211; UNIV MISSOURI, DEPT BIOCHEM/COLUMBIA//MO/65211; ACAD SCI CZECH REPUBL, INST EXPTL BOT/CR-16630 PRAGUE 6//CZECH REPUBLIC/; UNIV MISSOURI, DEPT AGRON/COLUMBIA//MO/65211

Journal: PLANT PHYSIOLOGY AND BIOCHEMISTRY, 1995, V33, N3 (MAY-JUN), P327-336

ISSN: 0981-9428

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number per ear at maturity by up to 30% and in some cases the total kernel weight per ear. The increase was due to a reduction in apical kernel abortion.

6/3,AB/8 (Item 5 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
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04047799 Genuine Article#: QK335 Number of References: 29
Title: INCREASE OF ENDOGENOUS ZEATIN RIBOSIDE BY INTRODUCTION OF THE
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(Abstract Available)

Author(s): GROOT SPC; BOUWER R; BUSSCHER M; LINDHOUT P; DONS HJ
Corporate Source: CTR PLANT BREEDING & REPROD RES, CPRO, DLO, DEPT DEV
BIOL, POB 16/6700 AA WAGENINGEN//NETHERLANDS/; CTR PLANT BREEDING &
REPROD RES, CPRO, DLO, DEPT VEGETABLE & FRUIT CROPS/6700 AA
WAGENINGEN//NETHERLANDS/

Journal: PLANT GROWTH REGULATION, 1995, V16, N1 (JAN), P27-36 ISSN: 0167-6903

Language: ENGLISH Document Type: ARTICLE

Abstract: We studied axillary meristem formation of the lateral suppressor (1s) mutant of tomato after elevating the endogenous cytokinin levels through introduction of the isopentenyltransferase (ipt) gene from Agrobacterium tumefaciens. Growth and development of several transformants were examined during in vitro culture. Transformants exhibited phenotypes varying in severity and were divided into four classes. A number of the ipt transformants had a normal phenotype, as non-transformed plants. Others showed a mild to severe 'cytokinin-like' phenotype. Transformants with a mild phenotype exhibited reduced internode length and reduced root development. Transformants with a severe phenotype showed even shorter internodes, loss of apical dominance, reduction of leaf size, production of callus at the basis of the shoots and absence of root development or development of green non-branching roots. The severity of the phenotype correlated well with the level of ipt gene expression, as measured by northern analysis. Transformants with a severe phenotype also exhibited increased levels of zeatin riboside, but zeatin levels were not elevated. The increase in endogenous zeatin riboside levels in the 1s mutant did not restore axillary meristem formation, but sometimes bulbous structures were formed in the initially 'empty' leaf axils. Several adventitious meristems and shoots developed from below the surface of these structures. It is concluded that a reduced level of cytokinins in the ls mutant shoots is not responsible for the absence of axillary meristem formation.

6/3,AB/9 (Item 6 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
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03927116 Genuine Article#: QT551 Number of References: 40
Title: THE EFFECT OF AUXIN ON CYTOKININ LEVELS AND METABOLISM IN
TRANSGENIC TOBACCO TISSUE EXPRESSING AN IPT GENE (Abstract Available)

Author(s): ZHANG R; ZHANG X; WANG J; LETHAM DS; MCKINNEY SA; HIGGINS TJV Corporate Source: AUSTRALIAN NATL UNIV, COOPERAT RES CTR PLANT SCI, POB 475/CANBERRA/ACT 2601/AUSTRALIA/; AUSTRALIAN NATL UNIV, COOPERAT RES CTR PLANT SCI/CANBERRA/ACT 2601/AUSTRALIA/; AUSTRALIAN NATL UNIV, RES SCH BIOL SCI, PLANT CELL BIOL GRP/CANBERRA/ACT 2601/AUSTRALIA/; CSIRO, DIV PLANT IND/CANBERRA/ACT 2601/AUSTRALIA/

Journal: PLANTA, 1995, V196, N1 (MAR), P84-94

ISSN: 0032-0935

Language: ENGLISH Document Type: ARTICLE

Abstract: The ipt gene from the T-DNA Agrobacterium tumefaciens was transferred to tobacco (Nicotiana tabacum L.) in order to study the control which auxin appears to exert over levels of cytokinin generated by expression of this gene. The transgenic tissues contained elevated levels of cytokinins, exhibited cytokinin and auxin autonomy and grew as shooty calli on hormone-free media. Addition of 1-naphthylacetic acid to this culture medium reduced the total level of cytokinins by 84% while 6-benzylaminopurine elevated the cytokinin level when added to media containing auxin. The cytokinins in the transgenic tissue were labelled with H-3 and auxin was found to promote conversion of zeatin-type cytokinins to H-3-labelled adenine derivatives. When the very rapid metabolism of exogenous [H-3]zeatin riboside was suppressed by a phenylurea derivative, a noncompetitive inhibitor of cytokinin oxidase, auxin promoted metabolism to adenine-type compounds. Since these results indicated that auxin promoted cytokinin oxidase activity in the transformed tissue, this enzyme was purified from the tobacco tissue cultures. Auxin did not increase the level of the enzyme per unit tissue protein, but did enhance the activity of the enzyme in vitro and promoted the activity of both glycosylated and non-glycosylated forms. This enhancement could contribute to the decrease in cytokinin level induced by auxin. Studies of cytokinin biosynthesis in the transgenic tissues indicated that transhydroxylation of isopentenyladenine-type cytokinins to yield zeatin-type cytokinins occurred principally at the nucleotide level.

6/3,AB/10 (Item 7 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
(c) 2002 Inst for Sci Info. All rts. reserv.

03826016 Genuine Article#: QH874 Number of References: 164
Title: MOLECULAR-GENETICS OF AUXIN AND CYTOKININ
Author(s): HOBBIE L; TIMPTE C; ESTELLE M
Corporate Source: INDIANA UNIV, DEPT BIOL/BLOOMINGTON//IN/47405; INDIANA UNIV, DEPT BIOL/BLOOMINGTON//IN/47405
Journal: PLANT MOLECULAR BIOLOGY, 1994, V26, N5 (DEC), P1499-1519
ISSN: 0167-4412

Language: ENGLISH Document Type: REVIEW

6/3,AB/11 (Item 8 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
(c) 2002 Inst for Sci Info. All rts. reserv.

03826015 Genuine Article#: QH874 Number of References: 113
Title: CYTOKININ METABOLISM - IMPLICATIONS FOR REGULATION OF
PLANT-GROWTH AND DEVELOPMENT

Author(s): BRZOBOHATY B; MOORE I; PALME K

Corporate Source: ACAD SCI CZECH REPUBL, INST BIOPHYS, KRALOVOPOLSKA 135/CR-61265 BRNO//CZECH REPUBLIC/; ACAD SCI CZECH REPUBL, INST BIOPHYS/CR-61265 BRNO//CZECH REPUBLIC/; UNIV OXFORD, DEPT PLANT SCI/OXFORD OX1 3RB//ENGLAND/

Journal: PLANT MOLECULAR BIOLOGY, 1994, V26, N5 (DEC), P1483-1497

ISSN: 0167-4412

Language: ENGLISH Document Type: REVIEW

6/3,AB/12 (Item 9 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
(c) 2002 Inst for Sci Info. All rts. reserv.

03731546 Genuine Article#: QB247 Number of References: 46

Title: STUNTED-PLANT-1, A GENE REQUIRED FOR EXPANSION IN RAPIDLY ELONGATING BUT NOT IN DIVIDING CELLS AND MEDIATING ROOT-GROWTH RESPONSES TO APPLIED CYTOKININ (Abstract Available)

Author(s): BASKIN TI; CORK A; WILLIAMSON RE; GORST JR

Corporate Source: UNIV MISSOURI, DIV BIOL SCI, 109 TUCKER

HALL/COLUMBIA//MO/65211; AUSTRALIAN NATL UNIV, RES SCH BIOL SCI, PLANT CELL BIOL GRP/CANBERRA/ACT 2601/AUSTRALIA/; UNIV TASMANIA, DEPT PLANT SCI/HOBART/TAS 7001/AUSTRALIA/

Journal: PLANT PHYSIOLOGY, 1995, V107, N1 (JAN), P233-243

ISSN: 0032-0889

Language: ENGLISH Document Type: ARTICLE

Abstract: To understand the control of spatial patterns of expansion, we have studied root growth in wild type and in the stunted plant 1 mutant, stp1, of Arabidopsis thaliana. We measured profiles of cell length and calculated the distribution of elongation rate. Slow growth of stp1 results both from a failure of dividing cell number to increase and from low elongation rates in the zone of rapid expansion. However, elongation of dividing cells was not greatly affected, and stp1 and wild-type callus grew at identical rates. Thus, rapid cellular expansion differs in mechanism from expansion in dividing cells and is facilitated by the STP1 gene. Additionally, there was no difference between stpl and wild-type roots for elongation in response to abscisic acid, auxin, ethylene, or gibberellic acid or for radial expansion in response to ethylene; however, stp1 responded to cytokinin much less than wild type. In contrast, both genotypes responded comparably to hormones when explants were cultured; in particular, there was no difference between genotypes in shoot regeneration in response to cytokinin. Thus, effects on root expansion mediated by cytokinin, but not effects mediated by other hormones or effects on other cytokinin-mediated responses, require the STP1 locus.

6/3,AB/13 (Item 10 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
(c) 2002 Inst for Sci Info. All rts. reserv.

02861394 Genuine Article#: MK413 Number of References: 26
Title: MORPHOMETRIC ANALYSIS OF THE GROWTH OF PHSP70-IPT TRANSGENIC
TOBACCO PLANTS (Abstract Available)

Author(s): VANLOVEN K; BEINSBERGER SEI; VALCKE RLM; VANONCKELEN HA; CLIJSTERS HMM

Corporate Source: LIMBURGS UNIV CENTRUM, DEPT SBG, UNIV CAMPUS/B-3610
DIEPENBEEK//BELGIUM/; LIMBURGS UNIV CENTRUM, DEPT SBG, UNIV CAMPUS/B-3610
DIEPENBEEK//BELGIUM/; UNIV INSTELLING ANTWERP, DEPT BIOL/B-2610
WILRIJK//BELGIUM/

Journal: JOURNAL OF EXPERIMENTAL BOTANY, 1993, V44, N268 (NOV), P 1671-1678

ISSN: 0022-0957

Language: ENGLISH Document Type: ARTICLE

Abstract: The effect of introducing a supplementary <code>ipt</code>-gene into the genome of Nicotiana tabacum L. cv. Petit Havana SR1 is studied on the morphological <code>plant</code> development. The <code>ipt</code>-gene, accounting for the biosynthesis of <code>cytokinins</code>, was coupled to the heat-inducible hsp70- promoter from Drosophila melanogaster. Besides the influence of the hormonal changes involved, the effects of the experimental conditions are examined, namely the in vitro growth conditions for selecting transformed <code>plants</code> and the heat treatment to induce <code>ipt</code>-gene expression.

The phenotype of the **plants** is determined by the tissue sensitivity to three factors: (1) heat treatment reduces stem elongation and diameter growth; (2) in vitro pre-cultivation also reduces stem elongation; and (3) expression of the **ipt**-gene stimulates diameter growth, induces debudding of the axillary shoots

and inhibits root development. In addition, axillary bud development indicates that in vitro cultivation affects ipt-gene expression.

6/3,AB/14 (Item 1 from file: 50)
DIALOG(R)File 50:CAB Abstracts
(c) 2002 CAB International. All rts. reserv.

02762279 CAB Accession Number: 931642994

Floral development and expression of floral homeotic genes are influenced by cytokinins.

Estruch, J. J.; Granell, A.; Hansen, G.; Prinsen, E.; Redig, P.; Onckelen, H. van; Schwarz-Sommer, Z.; Sommer, H.; Spena, A.

Max-Planck-Institut fur Zuchtungsforschung, Carl-von-Linne-Weg 10, 5000 Koln 30, Germany.

Plant Journal vol. 4 (2): p.379-384

Publication Year: 1993

ISSN: 0960-7412 --Language: English

Document Type: Journal article

Tobacco plants that are somatic mosaics for the expression of a cytokinin-synthesizing gene (isopentenyl transferase) have viviparous leaves and were obtained by inserting the maize transposon Ac into the untranslated leader sequence of the 35S-ipt gene. Epiphyllous buds can be either vegetative or floral. Floral adventitious buds can be either normal or abnormal. Abnormalities of floral development correlate with: (1) a local activation of the cytokinin-synthesizing gene; (2) a drastic increase in floral cytokinin content; and (3) a decrease in the steady-state levels of mRNA homologues of the homeotic genes DEFA, GLO and PLENA of Antirrhinum majus. Thus, these data show that cytokinins in planta are able to alter the development of floral organs and to decrease the expression of 3 homeotic floral genes. Nucleotide sequence data for the tobacco cDNA clone are deposited under EMBL Data Library accession number X67959. 29 ref.

6/3,AB/15 (Item 2 from file: 50)
DIALOG(R)File 50:CAB Abstracts
(c) 2002 CAB International. All rts. reserv.

02757131 CAB Accession Number: 930767659

Control of **cytokinin** levels by inhibitors of metabolism, symbiosis and genetic manipulation.

Hocart, C. H.; Letham, D. S.; Wang, J.; Cornish, E.; Parker, C. W. Research School of Biological Sciences, Australian National University, Canberra, ACT 2601, Australia.

Conference Title: Progress in plant growth regulation. Proceedings of the 14th international conference on plant growth substances, Amsterdam, 21-26 July, 1991

p.607-616

Publication Year: 1992

Editors: Karssen, C. M.; Loon, L. C. van; Vreugdenhil, D.

Publisher: Kluwer Academic Publishers -- Dordrecht, Netherlands

ISBN: 0-7923-1617-7 Language: English

Document Type: Conference paper

A review and discussion on increasing cytokinin levels either indirectly by inhibiting cytokinin N-glucosylation and alanine conjugation in radishes, maize, oats and soyabeans, or directly by increasing the level of cytokinins either by the incorporation of the cytokinin biosynthetic gene (ipt) into the plant genome to increase the production of N6-(isopent-2-enyl)adenosine-5'-phosp hate, or through the mediation of a symbiotic relationship, such as between a cytokinin -overproducing strain of Rhizobium and

pigeonpeas. 23 ref.

6/3,AB/16 (Item 3 from file: 50)
DIALOG(R)File 50:CAB Abstracts
(c) 2002 CAB International. All rts. reserv.

02708928 CAB Accession Number: 931638660

Viviparous leaves produced by somatic activation of an inactive cytokinin-synthesizing gene.

Estruch, J. J.; Prinsen, E.; Onckelen, H. van

MPI fur Zuchtungsforshchung, Carl-von-Linne Weg 10, W-5006 Koln 30, Germany.

Science (Washington) vol. 254 (5036): p.1364-1367

Publication Year: 1991

ISSN: 0036-8075 --Language: English

Document Type: Journal article

A chimaeric gene consisting of the CaMV 35S promoter and the isopentenyl transferase (ipt) gene of Agrobacterium tumefaciens, split by the Activator element of maize, was introduced into tobacco. Tobacco plants that are somatic mosaics for expression of a cytokinin-synthesizing ipt gene have viviparous leaves. Such a formation of shoots in an abnormal position represents a significant deviation from the usual organization of the plant body where a central axis produces shoots only in the axils of lateral leaf appendages and according to a precise phyllotactic pattern. This report links vivipary to the expression of a gene whose product is involved in the synthesis of the phytohormone cytokinin. 27 ref.

6/3,AB/17 (Item 4 from file: 50)
DIALOG(R)File 50:CAB Abstracts
(c) 2002 CAB International. All rts. reserv.

Alterations of endogenous cytokinins in transgenic plants using a chimeric isopentenyl transferase gene.

Medford, J. I.; Horgan, R.; El-Sawi, Z.; Klee, H. J.

Pl. Molec. Biol. Group, Monsanto Co., 700 Chesterfield Village Parkway, St. Louis, MO 63198, USA.

Plant Cell vol. 1 (4): p.403-413

Publication Year: 1989

ISSN: 1040-4651 --Language: English

Document Type: Journal article

Cytokinins appear to play an important role in the processes of plant development. The Agrobacterium tumefaciens isopentenyl transferase gene was placed under the control of a heat-inducible promoter (maize hsp70). The chimaeric gene was transferred to tobacco and Arabidopsis plants. Heat induction of transgenic plants caused isopentenyl transferase mRNA to accumulate and increased the level of zeatin 52-fold, zeatin riboside 23-fold and zeatin riboside 5'-monophosphate 2-fold. At the control temperature zeatin riboside and zeatin riboside 5'-monophosphate accumulated in transgenic plants to levels 3 and 7 times, respectively, over levels in wild-type plants. This uninduced cytokinin increase affected various aspects of development. In tobacco these effects included release of axillary buds, reduced stem and leaf area and an underdeveloped root system. In Arabidopsis reduction of root growth was also found. However, neither tobacco nor Arabidopsis transgenic plants showed any differences relative to wild-type plants in time of flowering. Unexpectedly, heat induction of cytokinins in transgenic plants produced no changes beyond those seen in the uninduced state. The lack of effect from

heat-induced increases could be a result of the transient increases in levels, direct cytokinin or indirect induction of negating factor(s), or lack of a corresponding level of competent cellular factors. Overall, the effects of the increased levels of endogenous cytokinins in non-heat-shocked transgenic plants seemed to be confined to aspects of growth rather than differentiation. Since no alterations in the programmed differentiation pattern were found with increased cytokinin levels, it is thought that this process may be controlled by components other than absolute cytokinin levels. 36 ref.

6/3,AB/18 (Item 1 from file: 357) DIALOG(R) File 357: Derwent Biotech Res (c) 2002 Thomson Derwent & ISI. All rts. reserv.

0199832 DBA Accession No.: 96-10012 MAT (Multi-Auto-Transformation) vector system 'marker-free transgenic plants can be visually selected by using the IPT gene as a positive marker - isopentenyl-transferase ipt gene selectable marker application in the visual selection of transgenic plant (conference abstract) AUTHOR: Ebinuma H; Sugita K; Matsunaga E; Yamakado M

CORPORATE AFFILIATE: Nippon-Paper

CORPORATE SOURCE: Central Research Laboratory, Nippon Paper Industries, Co., Ltd., 5-21-1, Oji, Kita-ku, Tokyo, Japan. email:LDW06374@niftyserve.or.jp

JOURNAL: Plant Physiol. (111, 2, Suppl., 42) 1996

ISSN: 0032-0889 CODEN: PLPHAY

CONFERENCE PROCEEDINGS: Plant Biology '96; 1996 Annual Meeting of the American Society of Plant Physiologists, San Antonio, TX, 27 July-2 August, 1996.

LANGUAGE: English

ABSTRACT: The ipt gene coding for isopentenyl-transferase which catalyzes cytokinin synthesis, was used as a positive marker to select transgenic plants. To remove the 35S-ipt gene from transgenic plants after transformation, the gene was combined into the maize (Zea mays) transposable element Ac the yeast site-specific-recombination system pSR1 or (pNPI132). This was termed a MAT (Multi-Auto-Transformation) vector system. Results indicated that the positive marker, the chimeric ipt gene, has a number of promising properties which make it an attractive alternative to the negative selectable marker genes. (1) Positive effects on cell division and differentiation of transgenic plants by cytokinin; (2) visual selection of transgenic plants with the chimeric ipt gene by morphological characteristics; (3) visual selection of marker-free transgenic plants without the chimeric ipt gene by morphological change. The MAT vector system enables the production of environmentally safe transgenic plants without sexual crosses and seed production, and pyramid multiple genes into a vegetatively propagated crop by repeated transformation. (0 ref)

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       S7
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                 S4 AND ZEA (W) MAYS
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 9/3, AB/1
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DIALOG(R) File
                5:Biosis Previews(R)
(c) 2002 BIOSIS. All rts. reserv.
09042188
           BIOSIS NO.: 199497050558
Cytokinins in plant pathogenic bacteria and developing cereal
  grains.
AUTHOR: Morris Roy O(a); Blevins Dale G; Dietrich Joseph T; Durley Richard
  C; Gelvin Stanton B; Gray John; Hommes Norman G; Kaminek Miroslav;
  Mathews Leslie J; et al
AUTHOR ADDRESS: (a) Dep. Biochem., Univ. Mo., Columbia, MO 65211**USA
JOURNAL: Australian Journal of Plant Physiology 20 (4-5):p621-637
1993
ISSN: 0310-7841
DOCUMENT TYPE: Article
RECORD TYPE: Abstract
LANGUAGE: English
ABSTRACT: Cytokinin analysis by immunoaffinity chromatography (IAC),
 high-performance liquid chromatography (HPLC) and radioimmunoassay (RIA)
 or enzyme-linked immunosorption assay (ELISA) has been used to study two
 separate topics: the role of tRNA in bacterial cytokinin
 biosynthesis and the changes in cytokinin concentration which occur
 during cereal grain development. Transfer RNA isopentenylation in the
 gall-forming plant pathogen Agrobacterium tumefaciens is encoded by
 the chromosomal miaA locus. Mutation of miaA reduces tRNA
 isopentenylation significantly and preliminary data suggest that turnover
 of isopentenylated tRNA is responsible for low level secretion of free
 N-6-isopentenyladenine (iP) by the bacteria. However, the major route of
 cytokinin biosynthesis by gall-forming plant pathogenic
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bacteria is not via tRNA turnover but by direct biosynthesis mediated by

dimethylallylpyrophosphate: 5'-AMP transferase (DMAPP:AMP transferase) encoded by such genes as <code>ipt</code>, tzs (from A. tumefaciens) or ptz (from Pseudomonas savastanoi). Analysis of <code>cytokinin</code> levels in developing wheat and rice grains in the period immediately following pollination showed large transient increases in zeatin (Z) and zeatin riboside (ZR) which coincided with the period of maximum endosperm cell division reported by others. Detailed analyses of <code>maize</code> kernels, where development can be staged readily, showed that Z and ZR concentrations peaked 9 days after pollination (DAP). During the period 8-10 DAP, <code>cytokinin</code> oxidase underwent a significant increase in specific activity, indicating that <code>cytokinin</code> catabolism was enhanced as endosperm cell division ended.

1993

9/3,AB/2 (Item 2 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
(c) 2002 BIOSIS. All rts. reserv.

06694909 BIOSIS NO.: 000088004327

ALTERATIONS OF ENDOGENOUS CYTOKININS IN TRANSGENIC PLANTS USING A CHIMERIC ISOPENTENYLTRANSFERASE GENE

AUTHOR: MEDFORD J I; HORGAN R; EL-SAWI Z; KLEE H J

AUTHOR ADDRESS: PLANT MOL. BIOL GROUP, MONSANTO CO., 700 CHESTERFIELD VILLAGE PARKWAY, ST. LOUIS, MO. 63198.

JOURNAL: PLANT CELL 1 (4). 1989. 403-414. 1989

FULL JOURNAL NAME: Plant Cell

CODEN: PLCEE

RECORD TYPE: Abstract

LANGUAGE: ENGLISH

ABSTRACT: Cytokins, a class of phytohormones, appear to play an important role in the processes of plant development. We genetically engineered the Argrobacterium tumerfaciens isopentenyl transferase gene, placing it under control of a heat-inducible promoter (maize hsp70). The chimeric hsp70 is isopentenyl transferase gene was transferred to tobacco and Arabidopsis plants. Heat induction of transgenic plants caused the isopentenyl transferase mRNA to accumulate and increase the level 52-fold, Zeatin riboside 23-fold, and Zeatin riboside 5'-monophosphate twofold. At the control temperature zeatin riboside and zeatin riboside 5'-monophosphate in transgenic plants accumula to levels 3 and 7 times, respectively, over levels in wild-type plants . This uninduced cytokinin increase affected various aspects of development. In tobacco, these effects included release axillary buds, reduced stem and leaf area, and an underdeveloped root system. In Arabidopsis, reduction of root growth was also found. However, neither tobacco nor Arabidopsis transgenic plants showed any differences relative to wild-type plants in time of flowering. Unexpectedly, heat induction of cytokinins in transgenic plants produced no changes beyond those seen in the uninduced state. The lack of effect from heat-induced increases could be a result of the transient increases in cytokinin levels, direct or indirect induction of negating factor(s), or lack of a corresponding level of competent cellular factors. Overall, the effects of the increased levels of endogenous cytokinins in non-heat shocked transgenic plants seemed to be confined to aspects of growth rather than differentiation. Since no alterations in the programmed differentiation pattern were found with increased cytokinin levels, this process may be controlled by components other than absolute cytokinin levels.

9/3,AB/3 (Item 1 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
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02767288 Genuine Article#: MC821 Number of References: 37
Title: THE ROLE OF CYTOKININ IN ORGANIZED DIFFERENTIATION OF VASCULAR
TISSUES (Abstract Available)

Author(s): ALONI R

Corporate Source: TEL AVIV UNIV, GEORGE S WISE FAC LIFE SCI, DEPT BOT/IL-69978 TEL AVIV//ISRAEL/

Journal: AUSTRALIAN JOURNAL OF PLANT PHYSIOLOGY, 1993, V20, N4-5, P 601-608

ISSN: 0310-7841

Language: ENGLISH Document Type: ARTICLE

Abstract: The role of cytokinin as a limiting and controlling factor in the differentiation of vascular tissues in the plant body is discussed. Cytokinin controls the early stages of fibre differentiation in Helianthus stems and the regeneration of vessels and sieve tubes around a wound in Coleus internodes. The influence of cytokinin on cell differentiation in the vascular tissues varies according to its physiological levels and the levels of auxin. Cytokinin induces an acropetal polar pattern of vessel regeneration around a wound in internodes of Coleus. Similarly, adventitious roots induce acropetal polar patterns of vessel maturation in hypocotyls of Cucurbita. Cytokinin increases the sensitivity of the vascular cambium to the auxin stimulation, resulting in the highest ratio of phloem/xylem under the optimal level of cytokinin. High levels of cytokinin promote callose production on sieve plates. Studies of transgenic plants with altered levels of cytokinin (overexpressing the ipt gene) confirm the involvement of cytokinin in vascular differentiation.

9/3,AB/4 (Item 2 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
(c) 2002 Inst for Sci Info. All rts. reserv.

02767276 Genuine Article#: MC821 Number of References: 87
Title: ALTERATIONS IN AUXIN AND CYTOKININ METABOLISM OF HIGHERPLANTS DUE TO EXPRESSION OF SPECIFIC GENES FROM PATHOGENIC
BACTERIA - A REVIEW (Abstract Available)

Author(s): HAMILL JD

Corporate Source: MONASH UNIV, DEPT GENET & DEV BIOL/CLAYTON/VIC 3168/AUSTRALIA/

Journal: AUSTRALIAN JOURNAL OF PLANT PHYSIOLOGY, 1993, V20, N4-5, P 405-423

ISSN: 0310-7841

Language: ENGLISH Document Type: ARTICLE

Abstract: This review deals with the physiological and morphological effects of altering the auxin/cytokinin balance in transgenic plants by expressing specific genes from pathogenic bacteria. Genes which have been used to alter auxin levels or sensitivity in transgenic plants include the iaaM/iaaH genes from Agrobacterium tumefaciens and A. rhizogenes; gene 5 and possibly gene 6b from A. tumefaciens; the rol B and possibly the rol A gene from A. rhizogenes and the iaaL gene from Pseudomonas syringae subsp. savastanoi (P. savastanoi). Genes which have been used to alter cytokinin levels in transgenic plants include the ipt gene from A. tumefaciens and the rol C gene from A. rhizogenes. A variety of biochemical mechanisms have been identified which result in alterations to phytohormone levels following expression of these genes in transgenic plants. Many of the effects on plant development are consistent with observations made following exogenous auxin and/or

cytokinin application to plant tissues, and the availability of these genes offers a new approach to the study of plant physiology using transformation methodology.

9/3,AB/5 (Item 3 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
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01117725 Genuine Article#: FX732 Number of References: 42
Title: DELAYED LEAF SENESCENCE IN TOBACCO PLANTS TRANSFORMED WITH
TMR, A GENE FOR CYTOKININ PRODUCTION IN AGROBACTERIUM (Abstract Available)

Author(s): SMART CM; SCOFIELD SR; BEVAN MW; DYER TA

Corporate Source: AFRC, INST GRASSLAND & ENVIRONM RES, WELSH PLANT BREEDING
STN, PLAS GOGERDDAN/ABERYSTWYTH SY23 3EB/DYFED/WALES/; JOHN INNES CTR
PLANT SCI RES, CAMBRIDGE LAB/NORWICH NR4 7UJ//ENGLAND/; JOHN INNES CTR
PLANT SCI RES, SAINSBURY LAB/NORWICH NR4 7UJ//ENGLAND/

Journal: PLANT CELL, **1991**, V3, N7, P647-656 Language: ENGLISH Document Type: ARTICLE

Abstract: The aim of this study was to investigate whether enhanced levels of endogenous cytokinins could influence plant development, particularly leaf senescence. Tobacco plants were transformed with the Agrobacterium tumefaciens gene tmr, under the control of the soybean heat shock promoter HS6871. This gene encodes the enzyme isopentenyl transferase, which catalyzes the initial step in cytokinin biosynthesis. After heat shock, the cytokinin level increased greatly and the level of tmr mRNA, undetectable at 20-degrees-C, rose and remained high for up to 8 hours. The levels of cytokinin and tmr mRNA were substantially lower by 24 hours. Transformed plants grown at 20-degrees-C were shorter, had larger side shoots, and remained green for longer than untransformed plants. The differences were more pronounced after several heat shocks of whole plants or defined areas of leaves. Our results demonstrated that plant morphology and leaf senescence can be manipulated by changing the endogenous level of cytokinins.

9/3,AB/6 (Item 4 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
(c) 2002 Inst for Sci Info. All rts. reserv.

00765104 Genuine Article#: EV082 Number of References: 38
Title: CYTOKININ CONTENT AND TISSUE DISTRIBUTION IN PLANTS
TRANSFORMED BY A RECONSTRUCTED ISOPENTENYL TRANSFERASE GENE
Author(s): SMIGOCKI AC

Corporate Source: USDA ARS, BELTSVILLE AGR RES CTR, PLANT MOLEC BIOL LAB/BELTSVILLE//MD/20705

Journal: PLANT MOLECULAR BIOLOGY, 1991, V16, N1, P105-115

Language: ENGLISH Document Type: ARTICLE

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09750490
           98226808
                      PMID: 9560269
  Agrobacterium transcriptional regulator Ros is a prokaryotic zinc finger
protein that regulates the plant oncogene ipt.
  Chou AY; Archdeacon J; Kado CI
  Davis Crown Gall Group, University of California, Davis, CA 95616, USA.
  Proceedings of the National Academy of Sciences of the United States of
America (UNITED
                 STATES)
                           Apr 28 1998, 95
                                                 (9)
                                                       p5293-8, ISSN
0027-8424
            Journal Code: PV3
  Contract/Grant No.: GM45550, GM, NIGMS
  Languages: ENGLISH
 Document type: Journal Article
 Record type: Completed
 Virulence genes of Agrobacterium tumefaciens are under the control of
positive and negative transcriptional regulators. We found that the
transcriptional regulator Ros controls expression of the plant
oncogene ipt, which encodes isopentenyl transferase, in
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A. tumefaciens. This enzyme is involved in biosynthesis of the plant growth hormone cytokinin in the host plant. An ipt promoter::cat reporter gene fusion showed a 10-fold increase in ipt promoter activity in A. tumefaciens ros mutant strains when compared with wild type. Also, increased levels (10- to 20-fold) of isopentenyl adenosine, the product of the reaction catalyzed by isopentenyl transferase , were detected in ros mutant strains. In vitro studies using purified Ros showed it binds directly to the ipt promoter. Analysis of the deduced Ros amino acid sequence identified a novel type of C2H2 zinc finger. In Ros the peptide loop spacing of the zinc finger is 9 amino acids as opposed to the invariant 12 amino acids in the classical C2H2 motif. Site-directed mutagenesis of Cys-82 and His-92 in this motif showed that these residues are essential for Zn2+ and DNA binding activities of Ros. The existence of such a regulator in Agrobacterium may be due to horizontal interkingdom retrotransfer of the ros gene from plant to bacteria.

12/3,AB/2 (Item 2 from file: 155) DIALOG(R)File 155:MEDLINE(R)

09587374 97446503 PMID: 9301091

Conditional transgenic **expression** of the **ipt** gene indicates a function for **cytokinins** in paracrine signaling in whole tobacco **plants**.

Faiss M; Zalubilova J; Strnad M; Schmulling T Universitat Tubingen, Lehrstuhl fur Allgemeine Genetik, Germany. Plant journal (ENGLAND) Aug 1997, 12 (2) p401-15, ISSN 0960-7412 Journal Code: BRU

Languages: ENGLISH

Document type: Journal Article

Record type: Completed

investigated study whether an increased production of the plant hormone cytokinin in roots, the main site of its synthesis and putative signaling organ, can influence developmental events, such as growth of axillary shoot meristems or leaf senescence, in the plant shoot. To this end, transgenic tobacco plants (Nicotiana tabacum L.) were generated that conditionally overproduce cytokinins. These plants harbour the ipt gene under the transcriptional control of a modified 35S promoter that is repressed in plants with high titers of tetracycline repressor protein. De-repression of transcription led to a rapid more than 50-fold increase of hormone concentration. The time course of changes in the steady-state levels of 16 different cytokinin metabolites, as a consequence of IPT enzyme activity, was monitored in different plant tissues. Zeatin riboside rst and most dramatically increased product; zeatin, and glucosides accumulated later. The consequences of the was first dihydrozeatin enhanced cytokinin synthesis remained mainly restricted to the site of hormone production. For example, de-repression of ipt gene transcription in lateral buds caused the growth of single buds only at the site of tetracycline application. In reciprocal grafts of transgenic plants with wild-type plants, no biological cytokinin effects, i.e. growth of lateral shoot meristems or sequential leaf senescence, were observed in the non-transgenic plant part. Also, the increase in steady-state levels of cytokinins remained restricted mainly to the transgenic part, despite a specific increase of the zeatin riboside concentration in the transpiration stream. These results question the role of **cytokinins** as a long-range root-to-shoot signal in correlative control of apical dominance and sequential leaf senescence of tobacco, and support the assumption that this hormone is relevant to paracrine signaling.

DIALOG(R) File 155: MEDLINE(R)

09302332 97245302 PMID: 9090061

Effects of seed-specific **expression** of a **cytokinin** biosynthetic gene on canola and tobacco phenotypes.

Roeckel P; Oancia T; Drevet J

Laboratoire associe Universite Blaise Pascal, INRA, Organisation et Variabilite des Genomes Vegetaux, Clermont-Ferrand, France.

Transgenic research (ENGLAND) Mar 1997, 6 (2) p133-41, ISSN 0962-8819 Journal Code: BRX

Languages: ENGLISH

Document type: Journal Article

Record type: Completed

The Agrobacterium tumefaciens isopentenyl transferase gene (ipt), a cytokinin biosynthetic gene, was placed under the control of 1.9 kb of promoter sequence from the 2S albumin AT2S1 gene isolated from an Arabidopsis thaliana library. The construct was introduced into canola (Brassica napus) and tobacco (Nicotiana tabacum). ipt were followed during embryo development of transgenic transcripts plants by northern hybridizations. The phenotype of transformed
plants from the T1 generation was analysed and we observed an increased branching of inflorescences in tobacco and canola plants expressing the ipt gene. Comparing with controls, the average
number of capsules and siliques in AT2S1-ipt plants was 82.6 and 24.8% higher, respectively. This result was correlated with an increase in cytokinin levels in transgenic plants, as revealed by RIA. Indeed, cytokinin contents of T1 AT2S1-ipt B. napus seeds were found 2.2-fold higher than cytokinin contents of control seeds, and AT2S1-ipt tabacum capsules contained Ν. 2.6-fold more cytokinins than control capsules. In tobacco, the average seed weight per capsule was lower in AT2S1-ipt plants while the seed number per silique and the average seed weight were not modified in canola carrying this construct. The average seed yield per plant was not significantly increased in AT2S1-ipt tobacco or canola plants.

12/3,AB/4 (Item 4 from file: 155) DIALOG(R)File 155:MEDLINE(R)

08865899 96165737 PMID: 8589740

Expression of the **isopentenyl transferase** gene is regulated by auxin in transgenic tobacco tissues.

Zhang XD; Letham DS; Zhang R; Higgins TJ

CSIRO Division of Plant Industry, Canberra, Australia.

Transgenic research (ENGLAND) Jan 1996, 5 (1) p57-65, ISSN 0962-8819 Journal Code: BRX

Languages: ENGLISH

Document type: Journal Article

Record type: Completed

The isopentenyl transferase gene (ipt) from Agrobacterium tumefaciens was isolated and introduced, via a disarmed binary vector, into tobacco using the Agrobacterium tumefaciens-mediated gene transfer system. The expression of the ipt gene was monitored by RNA hybridization, western blotting and cytokinin analysis. The addition of auxin to the media rapidly reduced the level of cytokinins in the transgenic tissues and this was associated with a reduction in IPT mRNA and protein levels. It is concluded that the hormone auxin can regulate expression of a gene involved in biosynthesis of the second hormone cytokinin. Although exogenous benzyladenine did not directly affect ipt gene expression, it did antagonize the effect of auxin on levels of cytokinins and IPT mRNA and protein.

12/3,AB/5 (Item 5 from file: 155) DIALOG(R)File 155:MEDLINE(R)

08765719 95195152 PMID: 7888614

Light-induced **expression** of **ipt** from Agrobacterium tumefaciens results in **cytokinin** accumulation and osmotic stress symptoms in transgenic tobacco.

Thomas JC; Smigocki AC; Bohnert HJ

Department of Biochemistry, University of Arizona, Tucson 85721.

Plant molecular biology (NETHERLANDS) Jan 1995, 27 (2) p225-35

ISSN 0167-4412 Journal Code: A60

Erratum in Plant Mol Biol 1995 Aug; 28(5) 965

Languages: ENGLISH

Document type: Journal Article

Record type: Completed

Cytokinins are plant growth regulators that induce shoot formation, inhibit senescence and root growth. Experiments with hydroponically grown tobacco plants, however, indicated that exogenously applied cytokinin led to the accumulation of proline and osmotin. These responses were also associated with environmental stress reactions, such as salt stress, in many plant species. To test whether increased endogenous cytokinin accumulation led to NaCl stress symptoms, the gene ipt from Agrobacterium tumefaciens, isopentenyl transferase , was transformed into encoding Nicotiana tabacum cv. SR-1 under the control of the light-inducible rbcS-3A promoter from pea. In high light (300 mumol PPFD m-2 s-1), ipt mRNA was detected and zeatin/zeatin glucoside levels were 10-fold higher than in control plants or when transformants were grown in low light (30 mumol PPFD m-2 s-1). High light treatment was accompanied by increased levels of proline and osmotin when compared to low light grown transformed untransformed control **plants**. Elevated in **planta** cytokinin levels induced responses also stimulated by salt stress, suggesting either common or overlapping signaling pathways are initiated independently by cytokinin and NaCl, setting in motion gene expression normally elicited by developmental processes such as flowering or environmental stress.

12/3,AB/6 (Item 6 from file: 155) DIALOG(R)File 155:MEDLINE(R)

08763598 96174437 PMID: 8592746

Inhibition of leaf senescence by autoregulated production of cytokinin.

Gan S; Amasino RM

Department of Biochemistry, University of Wisconsin, Madison, 53706-1569, USA.

Science (UNITED STATES) Dec 22 1995, 270 (5244) p1986-8, ISSN 0036-8075 Journal Code: UJ7

Languages: ENGLISH

Document type: Journal Article

Record type: Completed

Controlling expression of IPT, a gene encoding isopentenyltransferase (the enzyme that catalyzes the rate-limiting step in cytokinin biosynthesis), with a senescence-specific promoter results in the suppression of leaf senescence. Transgenic tobacco plants expressing this chimeric gene do not exhibit the developmental abnormalities usually associated with IPT expression because the system is autoregulatory. Because sufficient cytokinin is produced to retard senescence, the activity of the senescence-specific promoter is attenulated. Senescence-retarded leaves exhibit a prolonged, photosythetically active life-span. This result demonstrates that endogenously produced cytokinin can regulate senescence and provides a system to specifically manipulate the senescence program.

12/3,AB/7 (Item 7 from file: 155) DIALOG(R)File 155:MEDLINE(R)

08340865 95152559 PMID: 7849758

Promoter tagging with a promoterless **ipt** gene leads to **cytokinin**-induced phenotypic variability in transgenic tobacco **plants**:implications of gene dosage effects.

Hewelt A; Prinsen E; Schell J; Van Onckelen H; Schmulling T Universitat Tubingen, Lehrstuhl fur Allgemeine Genetik, Germany. Plant journal (ENGLAND) Dec **1994**, 6 (6) p879-91, ISSN 0960-7412 Journal Code: BRU

Languages: ENGLISH

Document type: Journal Article

Record type: Completed

Tobacco plants have been transformed with a T-DNA construct harboring a promoterless cytokinin-synthesizing ipt gene close to the right T-DNA border. Eighteen out of 85 transgenic clones displayed phenotypic alternations typical for an enhanced cytokinin production. Northern blot analysis confirmed the transcriptional activation of the introduced gene by tagged plant promoters. The concentration of cytokinins, expressed as zeatinriboside equivalents, was increased up to sevenfold in transgenic tissues. These increases in cytokinin levels resulted in major developmental changes. Transgenic clones exhibited to different levels traits of a general cytokinin -syndrome, i.e. reduced root growth, reduced apical dominance, reduced leaf surface, reduced growth of the stem and retarded leaf senescence or displayed localized and developmentally specific cytokinin-induced alterations in otherwise normally developing plants. These traits were in particular a simultaneous break of dormancy in all axillary buds before or at the onset of flowering or the reorientation of the developmental pathway of secondary meristems or terminally differentiated cells. This indicates that endogenously produced cytokinins not only influence different growth parameters but have the potential to alter differentiation pattern. The results show that stably inherited developmental alterations due to a general or localized cytokinin overproduction can be obtained by the promoter-tagging approach. The investigation of gene dosage effects in homozygote plants readdresses the question of threshold levels for cytokinin effects on the developmental program of plants.

12/3,AB/8 (Item 8 from file: 155) DIALOG(R)File 155:MEDLINE(R)

08011082 94035187 PMID: 8106083

Floral development and **expression** of floral homeotic genes are influenced by **cytokinins**.

Estruch JJ; Granell A; Hansen G; Prinsen E; Redig P; Van Onckelen H; Schwarz-Sommer Z; Sommer H; Spena A

Max-Planck-Institut fur Zuchtungsforschung, Koln, Germany.

Plant journal (ENGLAND) Aug 1993, 4 (2) p379-84, ISSN 0960-7412 Journal Code: BRU

Languages: ENGLISH

Document type: Journal Article

Record type: Completed

Tobacco plants that are somatic mosaics for the expression of a cytokinin -synthesizing gene have viviparous leaves. Epiphyllous buds can be either vegetative or floral. Floral adventitious buds can be either normal or abnormal. Abnormalities of floral development correlate with: (i) a local activation of the cytokinin-synthesizing gene, (ii) a drastic increase in floral cytokinin content, and (iii) a decrease in the steady-state levels of mRNA homologous of the homeotic genes DEFA,

GLO and PLENA of Antirrhinum majus. Thus, these data show in **planta** that **cytokinins**, a class of phytohormones, are able to alter the development of floral organs and to decrease the **expression** of three homeotic floral genes.

12/3,AB/9 (Item 9 from file: 155) DIALOG(R)File 155:MEDLINE(R)

07942495 94033311 PMID: 8219068

Cytokinin-mediated insect resistance in Nicotiana plants transformed with the ipt gene.

Smigocki A; Neal JW; McCanna I; Douglass L

Plant Molecular Biology Laboratory, U.S. Department of Agriculture, Beltsville, MD 20705.

Plant molecular biology (NETHERLANDS) Oct 1993, 23 (2) p325-35

ISSN 0167-4412 Journal Code: A60

Languages: ENGLISH

Document type: Journal Article

Record type: Completed

bacterial isopentenyl transferase (ipt) gene involved in cytokinin biosynthesis was fused with a promoter from the proteinase inhibitor II (PI-IIK) gene and introduced into Nicotiana plumbaginifolia. Transcripts of the ipt gene were wound-inducible in leaves of transgenic PI-II-ipt plants. In leaf disks excised from fully expanded leaves, transcript levels increased 25- to 35-fold within 24 h and by 48 h were reduced by about 50%. In flowering plants, message levels were 2- to 5-fold higher than in preflowering plants. These plants were used to test for defensive properties of cytokinins against insects. Manduca sexta larvae consumed up to 70% less of the PI-II-ipt leaf material on flowering plants than larvae feeding on controls. Normal development of Myzus persicae nymphs was also delayed. Approximately half as many nymphs reached adulthood on PI-II-ipt leaves than on controls. Zeatin and zeatinriboside levels in leaves remaining on PI-II-ipt plants after hornworm feeding were elevated by about 70-fold and the chlorophyll a/b content was double that of controls. Exogenous applications of zeatin to the PI-II-ipt leaves enhanced the level of resistance to the tobacco hornworm and almost completely inhibited normal development of the green peach aphid nymphs. Transcript levels of an acidic chitinase gene were low and minimally inducible in PI-II-ipt leaves. The mode of action of the cytokinin gene product on enhanced insect resistance is not clear but may involve the products of secondary metabolic pathways.

12/3,AB/10 (Item 10 from file: 155) DIALOG(R)File 155:MEDLINE(R)

07899388 93271451 PMID: 8499612

Regulatable endogenous production of **cytokinins** up to 'toxic' levels in transgenic **plants** and **plant** tissues.

Ainley WM; McNeil KJ; Hill J \overline{W} ; Lingle WL; Simpson RB; Brenner ML; Nagao RT; Key JL

Botany Department, University of Georgia, Athens 30602.

Plant molecular biology (NETHERLANDS) Apr 1993, 22 (1) p13-23, ISSN 0167-4412 Journal Code: A60

Contract/Grant No.: GM30317, GM, NIGMS

Languages: ENGLISH

Document type: Journal Article

Record type: Completed

The effects of **expressing** a chimeric gene consisting of a soybean heat shock gene promoter and a sequence that encodes an enzyme catalyzing the synthesis of a potent phytohormone, the **cytokinin** iPMP, have been analyzed in transgenic tobacco **plants**. The production of

endogenously produced several cytokinin effects previously undocumented. The differentiation of shoots independent of exogenous cytokinin from heat-treated transgenic plant leaf explants demonstrates that long-term heat treatments do not interfere with complex developmental processes. This extends the potential usefulness of heat shock gene promoters to conditionally express genes during windows of development that span several weeks.

12/3,AB/11 (Item 11 from file: 155) DIALOG(R) File 155: MEDLINE(R)

07642913 93012484 PMID: 1397692

Altered morphology in transgenic tobacco plants that overproduce cytokinins in specific tissues and organs.

Li Y; Hagen G; Guilfoyle TJ

Department of Biochemistry, University of Missouri, Columbia 65211. Developmental biology (UNITED STATES) Oct 1992, p386-95, ISSN 0012-1606 Journal Code: E7T

Languages: ENGLISH

Document type: Journal Article

Record type: Completed

An auxin-inducible bidirectional promoter from the soybean SAUR gene locus was fused to a reporter gene in one direction and a cytokinin biosynthetic gene in the opposite direction and the expression of these fused genes was examined in transgenic tobacco. The Escherichia coli uidA gene, which encodes the enzyme beta-glucuronidase (GUS), was used as the reporter gene and the Agrobacterium tumefaciens ipt gene, which encodes the enzyme isopentenyl transferase, was used as the cytokinin biosynthetic gene. These constructs allowed overproduction of cytokinins in tobacco in a tissue- and organ-specific manner. Localized overproduction of cytokinins was monitored using the GUS reporter gene and measured by an ELISA assay. The tissue- and organ-specific overproduction of cytokinins produced a number of morphological and physiological changes, including stunting, loss of apical dominance, reduction in root initiation and growth, either acceleration or prolonged delayed senescence in leaves depending on the growth conditions, adventitious shoot formation from unwounded leaf veins and petioles, altered nutrient distribution, and abnormal tissue development in stems. While some of these morphological changes result directly from the localized overproduction of cytokinins, other changes probably result from the mobilization of plant nutrients to tissues rich in cytokinins.

12/3, AB/12(Item 12 from file: 155) DIALOG(R)File 155:MEDLINE(R)

07580111 92192012 PMID: 1547783

Fasciation induction by the phytopathogen Rhodococcus fascians depends upon a linear plasmid encoding a cytokinin synthase gene.

Crespi M; Messens E; Caplan AB; van Montagu M; Desomer J Laboratorium voor Genetica, Universiteit Gent, Belgium.

journal (ENGLAND) **1992**, 11 Mar (3) p795-804, ISSN 0261-4189 Journal Code: EMB

Languages: ENGLISH

Document type: Journal Article

Record type: Completed

Rhodococcus fascians is a nocardiform bacteria that induces leafy galls (fasciation) on dicotyledonous and several monocotyledonous plants. The wild-type strain D188 contained a conjugative, 200 kb linear extrachromosomal element, pFiD188. Linear plasmid-cured strains were avirulent and reintroduction of this linear element restored virulence. Pulsed field electrophoresis indicated that the chromosome might also be a

linear molecule of 4 megabases. Three loci involved in phytopathogenicity have been identified by insertion mutagenesis of this Fi plasmid. Inactivation of the fas locus resulted in avirulent strains, whereas insertions in the two other loci affected the degree of virulence, yielding attenuated (att) and hypervirulent (hyp) bacteria. One of the genes within the fas locus encoded an isopentenyltranferase (IPT) with low homology to analogous proteins from Gram-negative phytopathogenic bacteria. IPT activity was detected after expression of this protein in Escherichia coli cells. In R.fascians, ipt expression could only be detected in bacteria induced with extracts from fasciated tissue. R.fascians strains without the linear plasmid but containing this fas locus alone could not provoke any phenotype on plants, indicating additional genes from the linear plasmid were also essential for virulence. These studies, the first genetic analysis of the interaction of a Gram-positive bacterium with plants, suggest that a novel mechanism for plant tumour induction has evolved in R.fascians independently from the other branches of the eubacteria.

12/3,AB/13 (Item 13 from file: 155) DIALOG(R)File 155:MEDLINE(R)

07445949 91363830 PMID: 1888890

Cytokinin content and tissue distribution in **plants** transformed by a reconstructed **isopentenyl transferase** gene. Smigocki AC

Plant Molecular Biology Laboratory, U.S. Department of Agriculture, Beltsville, MD 20705.

Plant molecular biology (NETHERLANDS) Jan 1991, 16 (1) p105-15, ISSN 0167-4412 Journal Code: A60

Languages: ENGLISH

Document type: Journal Article

Record type: Completed

The cytokinin gene, isopentenyl transferase (ipt), was placed under the control of a heat-inducible promoter from the melanogaster hsp70 gene and introduced into Nicotiana Drosophila plumbaginifolia by cocultivation with Agrobacterium Transformants were analyzed for organ-specific exp tumefaciens. expression, cytokinin levels and effects on plant development before and after the heat induction. The ipt gene transcripts were detected in leaves and stems but not roots of transgenic plants following a 2 hour, 45 degrees C treatment. Maximum mRNA levels observed occurred 2 hours after heat treatment and 46 hours later were detected only in leaves. Zeatin and zeatinriboside concentrations 2 hours after heat shock ranged from over 900 to 2000 pmol/g, representing a greater than 140- to 200-fold increase over uninduced levels. After 46 hours, approximately 50% of the cytokinins are still present in the leaves as opposed to much reduced levels in the stems. Transgenic plants were greener, shorter, had an underdeveloped root system, reduced leaf width, and increased growth of axillary buds. After a single heat treatment, **plants** exhibited a darker green pigment and continued growth of lateral buds. Transient accumulations of endogenous cytokinins following thermal induction did not appear to alter the plant 's preprogrammed pattern of differentiation.

12/3,AB/14 (Item 14 from file: 155) DIALOG(R)File 155:MEDLINE(R)

07333998 91355889 PMID: 2103461

Restoration of shooty morphology of a nontumorous mutant of Nicotiana glauca \times N. langsdorffii by **cytokinin** and the isopentenyltransferase gene.

Feng XH; Dube SK; Bottino PJ; Kung SD

Center for Agricultural Biotechnology, University of Maryland, College Park 20742.

Plant molecular biology (NETHERLANDS) Sep 1990, 15 (3) p407-20 ISSN 0167-4412 Journal Code: A60

Languages: ENGLISH

Document type: Journal Article

Record type: Completed

The shooty morphology of a nontumorous amphidiploid mutant of Nicotiana glauca Grah. x N. langsdorffii Weinm. was restored by cytokinins, whether exogenously applied or endogenously produced by transformation of the mutant with a transfer DNA (T-DNA) cytokinin-biosynthesis gene (isopentenyltransferase; ipt). Auxins alone did not confer this effect. Similar transformation was not achieved for the parental species. In the case of transformation with the ipt gene, selection of the transformed tissues was based on its hormone-independent growth in the presence of the antibiotic kanamycin. Transformed tissues exhibited a shooty morphology, indistinguishable from that of wildtype genetic tumors N. glauca x N. langsdorffii. This altered phenotype was caused by the presence and constitutive expression of the ipt gene. The insertion and expression of this gene in transformed tissues was confirmed by using the polymerase chain reaction (PCR) technique as well as conventional molecular hybridization analysis. Expression of the ipt gene led to an elevated level of cytokinin in the transformed mutant tissues. This evidence supports the notion that genetic tumors are caused, at least in part, by elevated levels of cytokinin in interspecific hybrids.

12/3,AB/15 (Item 15 from file: 155) DIALOG(R)File 155:MEDLINE(R)

05334767 90043783 PMID: 2811903

Transfer of the agrobacterial gene for **cytokinin** biosynthesis into tobacco **plants**]

Perenos v rasteniia tabaka agrobakterial'nogo gena biosinteza tsitokinina.

Iusibov VM; Pogosian GP; Andrianov VM; Piruzian ES

Molekuliarnaia genetika, mikrobiologiia i virusologiia (USSR) Jul 1989, (7) p11-3, ISSN 0208-0613 Journal Code: NMJ

Languages: RUSSIAN

Document type: Journal Article

Record type: Completed

The gene transfer into plants using the genetic engineering methods gives us the possibility to obtain transgeneric plants having acquired the new traits. Some bacterial genes can be used for this purpose. Obtaining of a transgeneric plant harbouring the cytokinin synthesis gene ipt (gene 4) from the T-DNA of Agrobacterium tumefaciens Ti-plasmid seems to be useful. The expression of tumor agrobacterial ipt gene in transformed plant cells interferes with the normal growth and regulation of the whole plant. The successful transfer of the cloned ipt gene from the recombinant plasmid pGV0319 into the tobacco plant using Agrobacterium vectors and succeeding regeneration of phenotypically normal transgenic plants are reported in the present paper.

12/3,AB/16 (Item 1 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
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12920094 BIOSIS NO.: 200100127243

Morphogenetic manifestations of the expression of the bacterial

ipt gene in regenerated tobacco plants in vitro.

AUTHOR: Makarova R V; Andrianov V M; Borisova T A; Piruzyan E S; Kefeli V I

JOURNAL: Fiziologiya Rastenii (Moscow) 44 (1):p11-19 January-February,

1997

MEDIUM: print ISSN: 0015-3303

DOCUMENT TYPE: Article RECORD TYPE: Abstract

LANGUAGE: Russian; Non-English SUMMARY LANGUAGE: English; Russian

ABSTRACT: The capacity of normal and transgenic tobacco plants (Nicotiana tabacum \hat{L} .) for regeneration, callus and organ formation after the expression of the active agrobacterial ipt gene was studied. In the cytokinin-transgenic explants (with the ipt gene), callus formation began only in the presence of 2,4-D and kinetin. Later the growth of callus tissue was hormone-independent. In the ipt regenerants, an abbreviated, particularly leafy shoot took shape, and the roots became implanted three to five days earlier. It was proposed that the morphological features of the ipt regenerants were conditioned by the specifics of their hormonal system.

1997

12/3,AB/17 (Item 2 from file: 5) DIALOG(R)File 5:Biosis Previews (R) (c) 2002 BIOSIS. All rts. reserv.

BIOSIS NO.: 200000409167

Transgenic Arabidopsis plants expressing ipt-gene.

AUTHOR: Werner T(a); Rupp H M; Scmuelling T; Van Onckelen H; Strnad M(a) AUTHOR ADDRESS: (a) Laboratory of Growth Regulators, Palacky University and Institute of Experimental Botany, Academy of Sciences of Czech Republic, 11 Slechtitelu, 783 71, Olomouc**Czech Republic

JOURNAL: Bulgarian Journal of Plant Physiology (Special Issue):p127 1998

MEDIUM: print

CONFERENCE/MEETING: 11th Congress of the Federation of European Societies

of Plant Physiology Varna, Bulgaria September 07-11, 1998 ISSN: 1310-4586

RECORD TYPE: Citation LANGUAGE: English

SUMMARY LANGUAGE: English

1998

12/3,AB/18 (Item 3 from file: 5) DIALOG(R)File 5:Biosis Previews(R) (c) 2002 BIOSIS. All rts. reserv.

BIOSIS NO.: 200000409160

Mechanisms controlling cytokinin levels in plant cells.

AUTHOR: Kaminek M(a); Motyka V; Gaudinova A; Dobrev P; Vankova R; Komanek D AUTHOR ADDRESS: (a) De Montfort University Norman Borlaug Institute for Plant Science, Institute of Experimental Botany, Academy of Sciences of the Czech Republic, Rozvojova 135, 16502, Prague 6**Czech Republic JOURNAL: Bulgarian Journal of Plant Physiology (Special Issue):p124 1998

MEDIUM: print

CONFERENCE/MEETING: 11th Congress of the Federation of European Societies

of Plant Physiology Varna, Bulgaria September 07-11, 1998

ISSN: 1310-4586 RECORD TYPE: Citation LANGUAGE: English

SUMMARY LANGUAGE: English

12/3,AB/19 (Item 4 from file: 5) DIALOG(R)File 5:Biosis Previews (R) (c) 2002 BIOSIS. All rts. reserv.

BIOSIS NO.: 199900073997

Autonomy to plant growth regulators and gene expression in

periwinkle cultures in vitro.

AUTHOR: Droual Anne-Marie(a); Hamdi Said(a); Creche Joel(a); Kevers Claire; Rideau Marc(a)

AUTHOR ADDRESS: (a) Lab. Biol. Mol. Biochim. Vegetale, EA 2106, Fac. Pharm., 31 Ave. Monge, F-37200 Tours-Cedex**France

JOURNAL: Journal of Plant Physiology 153 (5-6):p623-630 Nov., 1998

ISSN: 0176-1617

DOCUMENT TYPE: Article RECORD TYPE: Abstract LANGUAGE: English

ABSTRACT: To better understand the effect of habituation on gene expression in plant cells, we have compared the accumulation of specific mRNAs encoding respectively two proline-rich proteins, a chaperone protein and three enzymes linking primary and secondary metabolisms in two models of in vitro culture of periwinkle. These models consisted of two couples of a 2,4-dichlorophenoxyacetic acid-dependent/2,4-dichlorophenoxyacetic acid independent line in which autonomy to auxin and cytokinin was obtained either through habituation or through transformation with the isopentenyltransferase gene from Agrobacterium tumefaciens. Results showed that gene expression was modified by plant growth regulator autonomy but differently according to the type of autonomy: only the gene encoding a hydroxyproline-rich glycoprotein was regulated similarly in both PGR-independent lines. On the other hand, PGR autonomy did not lead to total insensitivity to exogenously-applied PGRs, and the two PGR autonomous lines did not accumulate indole alkaloids for different reasons.

1998

12/3,AB/20 (Item 5 from file: 5) DIALOG(R)File 5:Biosis Previews(R) (c) 2002 BIOSIS. All rts. reserv.

11666966 BIOSIS NO.: 199800448697

Expression of the bacterial ipt gene in Physcomitrella rescues mutations in budding and in plastid division.

AUTHOR: Reutter Kirsten; Atzorn Rainer; Hadeler Birgit; Schmuelling Thomas; Reski Ralf(a)

AUTHOR ADDRESS: (a) Albert-Ludwigs-Universitaet, Institut fuer Biologie II, Schaenzlestr. 1, D-79104 Freiburg**Germany

JOURNAL: Planta (Berlin) 206 (2):p196-203 Oct., 1998

ISSN: 0032-0935

DOCUMENT TYPE: Article RECORD TYPE: Abstract LANGUAGE: English

ABSTRACT: Development of Physcomitrella patens (Hedw.) B.S.G. starts with a filamentous protonema growing by apical cell division. As a developmental switch, some subapical cells produce three-faced apical cells, the so-called buds, which grow to form leafy shoots, the gametophores. Application of cytokinins enhances bud formation but no subsequent gametophore development in several mosses. We used the ipt gene of

Agrobacterium tumefaciens, encoding a protein which catalyzes the rate-limiting step in cytokinin biosynthesis, to transform two developmental Physcomitrella mutants. One mutant (P24) was defective in budding (bud) and thus did not produce three-faced cells, while the other one (PC22) was a double mutant, defective in plastid division (pdi), thus possessing at the most one giant chloroplast per cell, and in gametophore development (gad), resulting in malformed buds which could not differentiate into leafy gametophores. Expression of the ipt gene rescued the mutations in budding and in plastid division but not the one in gametophore development. By mutant rescue we provide evidence for a distinct physiological difference between externally applied and internally produced cytokinins. Levels of immunoreactive cytokinins and indole-3-acetic acid were determined in tissues and in culture media of the wild-type moss, both mutants and four of their stable ipt transformants. Isopentenyl-type cytokinins were the most abundant cytokinins in Physcomitrella, whereas zeatin-type cytokinins, the major native cytokinins of higher plants, were not detectable. Cytokinin as well as auxin levels were enhanced in ipt transgenics, demonstrating a cross-talk between both metabolic pathways. In all genotypes, most of the cytokinin and auxin was found extracellularly. These extracellular pools may be involved in hormone transport in the non-vascular mosses. We suggest that both mutants are defective in signal-transduction rather than in cytokinin metabolism.

1998

12/3,AB/21 (Item 6 from file: 5) DIALOG(R)File 5:Biosis Previews(R) (c) 2002 BIOSIS. All rts. reserv.

BIOSIS NO.: 199800404879 The cloning of rolC gene and over expression of cytokinins in Nicotiana tabacum.

AUTHOR: Jia Yan-Tao; Ma Mi(a); Qu Gui-Ping; Qian Zhong-Xing; Lin Zhong-Ping AUTHOR ADDRESS: (a) Inst. Bot., Chinese Acad. Sci., Beijing 100093**China JOURNAL: Acta Botanica Sinica 40 (3):p211-215 March, 1998

ISSN: 0577-7496

DOCUMENT TYPE: Article RECORD TYPE: Abstract

LANGUAGE: Chinese; Non-English SUMMARY LANGUAGE: Chinese; English

ABSTRACT: Using PCR method the rolC gene was amplified from Agrobacterium rhizogenes, and CaMV 35S/rolC expression vector pCaR was constructed. The chimeric gene via agrobacterium mediated procedure was transformed separately into the wild type tobacco (Nicotiana tabacum L. cv. W38) and the transgenic tobacco of ipt gene. The putative transgenic plants were assayed with Southern blot and RNA Dot blot analysis. The observation suggested that the transgenic tobacco exhibited the abnormal phenotypes as a consequence of the overproduction of cytokinins. Whereas the ELISA assay indicated that the cytokinins level increased separately in transgenic plants. The growth of the transgenic plants show multiple budding of shoots with short internodal length.

1998

12/3,AB/22 (Item 7 from file: 5) DIALOG(R)File 5:Biosis Previews(R) (c) 2002 BIOSIS. All rts. reserv.

11610186 BIOSIS NO.: 199800391950

Phenotypes of tobacco plants expressing genes for the synthesis of growth regulators.

AUTHOR: Hlinkova E(a); Obert B; Filipp D(a)

AUTHOR ADDRESS: (a) Dep. Genet., Fac. Nat. Sci., Comenius Univ., 84215

Bratislava**Slovakia

JOURNAL: Biologia Plantarum (Prague) 41 (1):p25-37 1998

ISSN: 0006-3134

DOCUMENT TYPE: Article RECORD TYPE: Abstract LANGUAGE: English

ABSTRACT: The expression of genes for synthesis of auxin (iaaM and iaaH) and cytokinins (ipt) was studied in tobacco plants transformed by two Agrobacterium tumefaciens strains C 58 and LBA 4404. The strain LBA 4404 carried binary vector plasmid pCB 1334 (ipt gene) and plasmid pCB 1349 (iaaM, iaaH and ila genes). Both plasmids carried reported gene for npt II. Obtained plants expressed incorporated genes. New proteins with molecular masses of about 74, 40, 26, 25, 21 and 17 kDa for wild plasmid pTi C58; 60, 36, 31.5, 27, 26 and 17 kDa for binary vector plasmid pCB 1334 and 74, 49, 36, 31.5, 26 and 25 kDa for binary vector plasmid pCB 1349 were found in the patterns of soluble proteins. Significant changes in the content of chlorophylls, especially chlorophyll a, were detected in the plants carrying ipt gene and in plants transformed by the wild strain C58 of A. tumefaciens. Tobacco plants expressing ipt gene and genes from T-DNA of pTi C58 plasmid were dwarf, and in comparison to the controls, they had thicker stems, and the surface of the leaf blades was reduced to 20-50%. Adventitious roots, growing from the stem, were typical for transformants overproducing auxins. Regenerants and transformants expressing genes from T-DNA of plasmid pTi C58 differed in the shape of the flowers and their fertility.

1998

12/3,AB/23 (Item 8 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
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11447051 BIOSIS NO.: 199800228383

Seed-specific expression of the isopentenyl transferase

gene (ipt) in transgenic tobacco.

AUTHOR: Ma Qing-Hu; Zhang Ren; Hocart Charles H(a); Letham David S; Higgins Thomas J V

AUTHOR ADDRESS: (a) Plant Biol. Group, Research School Biol. Sciences, Australian National Univ., GPO Box 475, Canbe**Australia

JOURNAL: Australian Journal of Plant Physiology 25 (1):p53-59 1998

ISSN: 0310-7841

DOCUMENT TYPE: Article RECORD TYPE: Abstract LANGUAGE: English

ABSTRACT: The Agrobacterium tumefaciens gene encoding <code>isopentenyl</code> <code>transferase</code> (<code>ipt</code>), a <code>cytokinin</code> biosynthetic gene, was fused to a promoter from a seed-specific gene, vicilin, and introduced into tobacco cells. Intact fertile <code>plants</code> were generated. The <code>expression</code> of the vicilin-<code>ipt</code> gene was shown to be confined to seed and resulted in enhanced levels of <code>cytokinins</code> in the developing seeds and increased seed protein content. Using a simplified quantification method, a significant increase in the levels of endogenous <code>cytokinins</code> was recorded at 16-21 days after flowering. The growth of the transgenic <code>plants</code> and the development of the seeds appeared to be normal.

12/3,AB/24 (Item 9 from file: 5) DIALOG(R)File 5:Biosis Previews(R) (c) 2002 BIOSIS. All rts. reserv.

BIOSIS NO.: 199800218198 11436866

Controlled cytokinin production in transgenic tobacco using a copper-inducible promoter.

AUTHOR: McKenzie Marian Jane(a); Mett Vadim; Reynolds Paul Hugh Stewart; Jameson Paula Elizabeth

AUTHOR ADDRESS: (a) Dep. Plant Biol. Biotechnol., Massey Univ., Private Bag 11222, Palmerston North**New Zealand

JOURNAL: Plant Physiology (Rockville) 116 (3):p969-977 March, 1998

ISSN: 0032-0889

DOCUMENT TYPE: Article RECORD TYPE: Abstract LANGUAGE: English

ABSTRACT: The cytokinin group of plant hormones regulates aspects of plant growth and development, including the release of lateral buds from apical dominance and the delay of senescence. In this work the native promoter of a cytokinin synthase gene (ipt) was removed and replaced with a Cu-controllable promoter. Tobacco (Nicotiana tabacum L. cv tabacum) transformed with this Cu-inducible ipt gene (Cu-ipt) was morphologically identical to controls under noninductive conditions in almost all lines produced. However, three lines grew in an altered state, which is indicative of cytokinin overproduction and was confirmed by a full cytokinin analysis of one of these lines. The in vitro treatment of morphologically normal Cu-ipt transformants with Cu2+ resulted in delayed leaf senescence and an increase in cytokinin concentration in the one line analyzed. In vivo, inductive conditions resulted in a significant release of lateral buds from apical dominance. The morphological changes seen during these experiments may reflect the spatial aspect of control exerted by this gene expression system, namely expression from the root tissue only. These results confirmed that endogenous cytokinin concentrations in tobacco transformants can be temporally and spatially controlled by the induction of ipt gene expression through the Cu-controllable geneexpression system.

1998

12/3,AB/25 (Item 10 from file: 5) DIALOG(R)File 5:Biosis Previews(R) (c) 2002 BIOSIS. All rts. reserv.

BIOSIS NO.: 199800053168

Studies of cytokinin action and metabolism using tobacco plants expressing either the ipt or the GUS gene controlled by a chalcone synthase promoter. II. ipt and GUS gene expression, cytokinin levels and metabolism.

AUTHOR: Wang Jian; Letham D S(a); Cornish Edwina; Wei K; Hocart C H; Michael M; Stevenson K R

AUTHOR ADDRESS: (a) Cooperative Res. Centre Plant Sci., Res. Sch. Biological Sci., Australian Natl. Univ., GPO Box 4**Australia

JOURNAL: Australian Journal of Plant Physiology 24 (5):p673-683 1997

ISSN: 0310-7841

DOCUMENT TYPE: Article RECORD TYPE: Abstract

LANGUAGE: English

ABSTRACT: The expression of GUS and ipt genes under control of a chalcone synthase (chs) promoter (PCHS) has been determined in tobacco (Nicotiana tabacum L.) plants and related to the development of plants expressing the chimaeric PCHS-ipt gene. GUS gene expression, which served as a model for the expression of the ipt gene, was highest in the internal phloem tissue of stems, in mature leaf laminae and in the upper part of corollas when fully open. Expression of the PCHS-ipt gene was assessed by quantifying the cytokinins produced, by determining incorporation of (3H) adenine into cytokinins and by quantifying ipt mRNA. Results from these studies were in general agreement with those based on expression of the PCHS-GUS gene. The chs promoter controlled expression of the ipt gene with some degree of tissue and temporal specificity. Expression of the ipt gene markedly elevated the cytokinin level in mature leaf laminae and the upper stems of flowering plants. The former was associated with retardation of leaf senescence and increased rates of transpiration due to changes in number, size and aperture of stomata, while the latter was associated with development of lateral shoots. In shoot tip cultures, 2-fold elevations in endogenous cytokinin level caused clear changes in development and this is discussed in relation to current concepts concerning the hormonal control of plant development. Using the transgenic tobacco tissues, it was shown that cis-zeatin is a substrate for cytokinin oxidase, that cis-zeatin is not converted to trans-zeatin in these tissues and that the endogenous cytokinin level influences the level of cytokinin oxidase activity in tissue and the rate of degradation of exogenous zeatin riboside to adenosine. 1997

12/3,AB/26 (Item 11 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
(c) 2002 BIOSIS. All rts. reserv.

11271835 BIOSIS NO.: 199800053167
Studies of cytokinin action and metabolism using tobacco plants
expressing either the ipt or the GUS gene controlled by a
chalcone synthase promoter. I. Developmental features of the transgenic
plants.

AUTHOR: Wang Jian; Letham D S(a); Cornish Edwina; Stevenson K R AUTHOR ADDRESS: (a) Cooperative Res. Centre Plant Sci., Res. Sch. Biological Sci., Australian Natl. Univ., GPO Box 4**Australia JOURNAL: Australian Journal of Plant Physiology 24 (5):p661-672 1997 ISSN: 0310-7841

DOCUMENT TYPE: Article RECORD TYPE: Abstract LANGUAGE: English

ABSTRACT: A chimaeric cytokinin biosynthetic gene was constructed by placing the coding region of the bacterial ipt gene under the control of a chalcone synthase (chs) promoter (PCHS) from Antirrhinum majus. The PCHS-ipt gene was transferred to tobacco (Nicotiana tabacum L.). To provide control plants for studies of the effect of expression of this gene on plant development, a PCHS beta-glucuronidase gene fusion was also introduced into tobacco. Expression of the PCHS-ipt gene caused release of axillary buds, inhibition of root development, retardation of leaf senescence, elevation of chlorophyll levels, delay in onset of flowering and retardation of flower development. These effects, which were quantified in PCHS-ipt plants, have previously been associated with expression of ipt genes controlled by heat shock or other promoters. Additional effects of ipt gene

expression characterized in PCHS-ipt plants included growth of leafy shoots from the primary root, change in leaf shape with the production of broader and larger leaves, induction of expansion of excised leaf discs and development of leaves with an enlarged midrib and enlarged veins. A particularly striking effect of the expression of the PCHS-ipt gene was development of thicker stems due mainly to increase of pith tissue caused by an enhancement of both cell division and cell enlargement. Node number per primary stem was also increased. Endogenous cytokinin and applied auxin interacted antagonistically to affect both root and stem development in plants cultured in vitro. The leaves of PCHS-ipt transformed plants exhibited increased transpiration rates and reduced diffusion resistance associated with increased number of stomata and modified stomatal dimensions. The above changes, which were associated with elevated endogenous cytokinin levels, are discussed in relation to previous studies with ipt gene transformed plants and to some aspects of normal plant development.

1997

12/3,AB/27 (Item 12 from file: 5) DIALOG(R)File 5:Biosis Previews (R) (c) 2002 BIOSIS. All rts. reserv. 11060001 BIOSIS NO.: 199799681146 Taproot-specific expression of a cytokinin biosynthesis gene (ipt) in transgenic sugarbeet. AUTHOR: Smigocki Ann C(a); McCanna Iris J; Ivic Snezana; Snyder Gordon W; Sicher Richard C; Owens Lowell D AUTHOR ADDRESS: (a) USDA-ARS Plant Mol. Biol. Lab., Beltsville, MD 20705** USA JOURNAL: Plant Physiology (Rockville) 114 (3 SUPPL.):p303 1997 CONFERENCE/MEETING: PLANT BIOLOGY '97: 1997 Annual Meetings of the American Society of Plant Physiologists and the Canadian Society of Plant Physiologists, Japanese Society of Plant Physiologists and the Australian Society of Plant Physiologists Vancouver, British Columbia, Canada August 2-6, 1997 ISSN: 0032-0889 RECORD TYPE: Citation LANGUAGE: English 1997

12/3,AB/28 (Item 13 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
(c) 2002 BIOSIS. All rts. reserv.

10897302 BIOSIS NO.: 199799518447

The role of cytokinin biosynthetic gene in regulating the expression of a class of pathogenesis-related protein genes in tobacco plants.

AUTHOR: Ma Qing-Hu Song Yan-Ru; Sun Jing-San

AUTHOR ADDRESS: Inst. Bot., Acad. Sinica, Beijing 100093**China JOURNAL: Acta Botanica Sinica 38 (11):p870-874 1996

ISSN: 0577-7496

RECORD TYPE: Abstract

LANGUAGE: English

SUMMARY LANGUAGE: English; Chinese

ABSTRACT: The **expression** characteristics of a class of pathogenesis-related protein (PR) genes, namely basic chitinase, beta-1, 3-glucanase, osmotin and extensin. were studied in tobacco (Nicotiana tabacum cv. Wisconsin 38) **plants**. RNA blot hybridization showed

that these four genes were regulated in a developmental and organ-specific manner in tobacco. In the transgenic fascicular shoots which contained the active **cytokinin** biosynthetic gene (**ipt** gene) from Agrobacterium tumefaciens, the **expressions** of these four genes were co-regulated by overproduction of endogenous **cytokinins** and vector effect. **Cytokinins** reduced the **expressions** while vector effect induced the **expressions** of these four genes. Heat shock also decreased the steady-state levels of the four RNAs. These data suggest a complex regulation of PR genes.

1996

12/3,AB/29 (Item 14 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
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10887646 BIOSIS NO.: 199799508791
Auxin-cytokinin interactions in wild-type and transgenic tobacco.
AUTHOR: Eklof Staffan(a); Astot Crister(a); Blackwell John(a); Moritz Thomas(a); Olsson Olof; Sandberg Goran(a)
AUTHOR ADDRESS: (a)Dep. Forest Genetics and Plant Physiol., Swed. Univ. Agric. Sci., S-901 83 Umea**Sweden
JOURNAL: Plant and Cell Physiology 38 (3):p225-235 1997
ISSN: 0032-0781
RECORD TYPE: Abstract
LANGUAGE: English

ABSTRACT: Cytokinins and auxins are important regulators of plant growth and development, but there is incomplete and conflicting evidence that auxins affect cytokinin metabolism and vice versa. We have investigated these interactions in Nicotiana tabacum L. by separate in planta manipulation of levels of the hormones followed by analysis of the induced changes in the metabolism of the other hormone. Cytokinin-overproducing plants (expressing the Agrobacterium tumefaciens ipt gene) had lower than wild-type levels of free IAA, and reduced rates of IAA synthesis and turnover, but there were no differences in the profiles of metabolites they produced from fed IAA. Similarly, auxin-overproducing plants (expressing the A. tumefaciens iaaM and iaaH genes), had lower levels of the major cytokinins than wild-type plants and lower cytokinin oxidase activity, but there were no differences in the profiles of metabolites they produced from fed cytokinins. The data demonstrate that cytokinin or auxin overproduction decreases the content of the other hormone, apparently by decreasing its rate of synthesis and/or transport, rather than by increasing rates of turnover or conjugation. Implications for the importance of cytokinin:auxin ratios in plant development are considered.

1997

12/3,AB/30 (Item 15 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
(c) 2002 BIOSIS. All rts. reserv.

10807972 BIOSIS NO.: 199799429117

Molecular analysis of the virulence determinants of the phytopathogen Rhodococcus fascians.

AUTHOR: Vereecke Danny; Temmerman Wim; Maes Tania; Van Montagu Marc;

AUTHOR ADDRESS: Lab. Genetica, Dep. Genet., Flanders Interuniversity Inst. Biotechnol., Univ. Gent, K.L. Ledeganckst**Belgium

JOURNAL: Mededelingen Faculteit Landbouwkundige en Toegepaste Biologische

Wetenschappen Universiteit Gent 61 (2A):p231-240 1996 RECORD TYPE: Abstract LANGUAGE: English

ABSTRACT: Most of the virulence determinants of the Gram-positive phytopathogen Rhodococcus fascians reside on a 200-kb linear plasmid, pFiD188. Three major groups of virulence genes can be distinguished on pFiD188. The fas locus contains six genes that encode proteins for cytochrome P450-linked electron transport and an isopentenyl transferase, the actions of which ultimately result in the formation of a hypermodified cytokinin. The att locus is a complex region of 20-kb that encompasses arginine biosynthetic genes and several genes related to polyketide biosynthesis; it probably encodes proteins for the formation of a complex molecule that would enhance susceptibility to infection or sensibility to suboptimal amounts of cytokinins. The third group of virulence genes contains regulatory proteins that are involved in control of the expression of fas and art both on the transcriptional and translational level. Besides these pFiD188-encoded loci one chromosomal virulence locus vic, was characterized. It is possibly involved in the catabolism of a gall specific compound that functions as a specialized carbon and nitrogen source for R. fascians.

1996

12/3,AB/31 (Item 16 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
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10788777 BIOSIS NO.: 199799409922
Regulation of cytokinin oxidase activity in tobacco callus expressing the T-DNA ipt gene.

AUTHOR: Redig Pascale; Motyka Vaclav; Van Onckelen Henri A; Kaminek Miroslav(a)

AUTHOR ADDRESS: (a) De Montfort Univ. Norman Borlaug Centre Plant Sci., Inst. Exp. Bot., Acad. Sci. Czech Republic, **Czech Republic

JOURNAL: Physiologia Plantarum 99 (1):p89-96 1997

ISSN: 0031-9317 RECORD TYPE: Abstract LANGUAGE: English

ABSTRACT: There are indications that the cytokinin content in transgenic tissues expressing the cytokinin biosynthetic ipt gene is under metabolic control, which prevents the accumulation of cytokinins to lethal levels. The objective of this study was to investigate the relationships between the content of endogenous cytokinins and the activity of cytokinin oxidase (which is believed to be a copper-containing amine oxidase, EC 1.4.3.6.) in ipt transgenic tobacco callus. In addition, the effect of exogenously applied N-6-benzyladenine (BA) on this relationship was examined. Endogenous cytokinin concentrations were measured in callus of Nicotiana tabacum L. cv. Petit Havana SR1 transformed with the ipt of Agrobacterium tumefaciens under the control of a light-inducible promoter and in non-transformed tissue using LC-tandem mass spectrometry. The activity of cytokinin oxidase was estimated by measuring the conversion of (2,8-3H)N-6-(DELTA-2-isopentenyl)adenine to (3H) adenine by enzyme preparations in vitro. The 14-day-old ipt -transformed callus contained a 25-fold higher amount of cytokinins as compared to the non-transformed tissue. Mainly zeatin- and dihydrozeatin types of cytokinins (free bases, ribosides, nucleotides and O-glucosides) accumulated in the ipt transgenic tissue. The cytokinin pool of both ipt-transformed and non-transformed tissues consisted predominantly of cytokinins that are either resistant to cytokinin oxidase attack (nucleotides and

O-glucosides of cytokinins and cytokinins bearing N-6-saturated side chain) or have a low affinity for the enzyme (zeatin and its riboside). The former represented 71.6 and 74.8% and the latter 27.7 and 24.4% of the pool of endogenous cytokinins in ipt -transformed and non-transformed tissues, respectively. Enzyme preparations from ipt-transformed tissue exhibited 1.5-fold higher cytokinin oxidase activity compared with that observed in control tissues. Application of exogenous BA affected the total levels of cytokinins of the two tissue fines in different ways. The cytokinin content increased by 1.7- and 1.5-fold in ipt -transformed tissues 6 and 12 h after BA application, respectively, while it declined in the non-transformed control by 1.6- to 2.0-fold between 3 and 12 h after BA application. The increase in cytokinin content in the ipt callus is due to an increase of zeatin- and dihydrozeatin-type cytokinins (nucleotides, ribosides and free bases) leading to an enhanced accumulation of O-glucosides after 12 h. Following BA treatment, the cytokinin oxidase activity increased up to 1.8-fold in ipt-transformed and 1.6-fold in nontransformed tissues. The levels of isopentenyl-type cytokinins were near the detection limit; however, the enhancement of cytokinin oxidase activity after BA treatment in both tissue lines was correlated with the content of preferred substrate of the enzyme, N-6-(DELTA-2-isopentenyl)adenosine.

1997

12/3,AB/32 (Item 17 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
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10749434 BIOSIS NO.: 199799370579
Combined effects of auxin transport inhibitors and cytokinin:
 Alterations of organ development in tobacco.
AUTHOR: Strabala Timothy J; Wu Yan H; Li Yi(a)
AUTHOR ADDRESS: (a)Division Biol., Kansas State Univ., Manhattan, KS 66506-4901**USA
JOURNAL: Plant and Cell Physiology 37 (8):p1177-1182 1996
ISSN: 0032-0781
RECORD TYPE: Abstract
LANGUAGE: English

ABSTRACT: We have examined the effects of the auxin transport inhibitors 1-naphthylphthalamic acid (NPA) and 2,3,5-triiodobenzoic acid (TIBA) on leaf morphogenesis of transgenic Nicotiana tabacum (cv. Xanthi) plants expressing the Agrobacterium tumefaciens cytokinin biosynthetic gene, ipt. We have observed the formation of saucer-shaped leaf-like organs at the shoot apex and at lateral buds. The formation of apical saucer-shaped leaf-like organs can be duplicated by the application of exogenous NPA and cytokinin to wild-type tobacco seedlings. We have also observed adventitious leaf-like organs with altered petiole and blade morphology in the transgenic plants treated with auxin transport inhibitors. These results suggest that the combination of diminished auxin transport and elevated cytokinin can lead to alterations in leaf development in tobacco.

1996

12/3,AB/33 (Item 18 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
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10697808 BIOSIS NO.: 199799318953

Changes in cytokinin content and cytokinin oxidase activity in response to derepression of ipt gene transcription in transgenic tobacco calli and plants.

AUTHOR: Motyka Vaclav; Faiss Martin; Strnad Miroslav; Kaminek Miroslav; Schmuelling Thomas(a)

AUTHOR ADDRESS: (a)Universitaet Tuebingen, Lehrstuhl fuer Allgemeine Genetik, Auf der Morgenstelle 28, D-72076 Tueb**Germany

JOURNAL: Plant Physiology (Rockville) 112 (3):p1035-1043 1996

ISSN: 0032-0889

RECORD TYPE: Abstract LANGUAGE: English

ABSTRACT: Metabolic control of cytokinin oxidase by its substrate was investigated in planta using wild-type (WT) and conditionally ipt gene-expressing transgenic (IPT) tobacco (Nicotiana tabacum L.) callus cultures and plants. The derepression of the tetracycline (Tc)-dependent ipt gene transcription was followed by a progressive, more than 100-fold increase in total cytokinin content in IPT calli. The activity of cytokinin oxidase extracted from these calli began to increase 16 to 20 h after gene derepression, and after 13 d it was 10-fold higher than from Tc-treated WT calli. An increase in cytokinin oxidase activity, as a consequence of elevated cytokinin levels, was found in detached leaves (8-fold after 4 d) and in roots of intact plants (4-fold after 3 d). The partially purified cytokinin oxidase from WT, repressed IPT, and Tc-derepressed IPT tobacco calli exhibited similar characteristics. it had the same broad pH optimum (pH 6.5-8.5), its activity in vitro was enhanced 4-fold in the presence of copper-imidazole, and the apparent K-m(N-6-(DELTA-2isopentenylladenine) values were in the range of 3.1 to 4.9 mu-M. The increase in cytokinin oxidase activity in cytokinin-overproducing tissue was associated with the accumulation of a glycosylated form of the enzyme. The present data indicate the substrate induction of cytokinin oxidase activity in different tobacco tissues, which may contribute to hormone homeostasis.

1996

12/3,AB/34 (Item 19 from file: 5) DIALOG(R)File 5:Biosis Previews(R) (c) 2002 BIOSIS. All rts. reserv.

BIOSIS NO.: 199699260488 10639343

Cytokinin metabolites and gradients in wild type and transgenic tobacco with moderate cytokinin over-production.

AUTHOR: Ekloff Staffan; Astot Crister; Moritz Thomas; Blackwell John; Olsson Olof; Sandberg Goran(a)

AUTHOR ADDRESS: (a) Dep. Forest Genetics Plant Physiol., Swed. Univ. Agric. Sch., S-901 85 Umea**Switzerland

JOURNAL: Physiologia Plantarum 98 (2):p333-344 1996

ISSN: 0031-9317

DOCUMENT TYPE: Article RECORD TYPE: Abstract LANGUAGE: English

ABSTRACT: A binary T-DNA plant expression vector carrying a promoterless isopentenyl transferase (ipt) gene was constructed and used to transform Nicotiana tabacum L. Several primary transformants were obtained that displayed a range of phenotypes characteristic of cytokinin over-production. Two of the transformants with moderately altered phenotypes, both of which produced viable offspring and expressed the ipt gene at a low level, were selected for use in studies of the regulation of cytokinin

metabolism. Both lines were found to contain high concentrations of zeatin-7-glucoside (Z7G), indicating that Z7G can accumulate in plants even when the rate of endogenous overproduction of cytokinins is low. This supports the hypothesis that 7-glucosidation is an important step in the regulation of zeatin (Z) levels. Very sharp gradients in concentration of cytokinin riboside and ribotides, related to age of tissue and distance from the apex, were found in both wild type and transformed plants, which could be important in developmental regulation and could also account for some of the discrepancies between reported cytokinin levels in various plants. Intriguingly, however, although the combined level of zeatin riboside and ribotide was much higher in the transformed plants than in wild type, the combined level of isopentenyl riboside and ribotide was lower.

1996

12/3,AB/35 (Item 20 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
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10616456 BIOSIS NO.: 199699237601
Analysis of cytokinin metabolism in ipt transgenic tobacco by liquid chromatography-tandem mass spectrometry.
AUTHOR: Redig Pascale(a); Schmuelling Thomas; Van Onckelen Harry AUTHOR ADDRESS: (a)Univ. Antwerp, Dep. Biol., Universiteitsplein 1, B-2610 Antwerpen, Belgium**Germany
JOURNAL: Plant Physiology (Rockville) 112 (1):p141-148 1996
ISSN: 0032-0889

DOCUMENT TYPE: Article RECORD TYPE: Abstract LANGUAGE: English

ABSTRACT: The endogenous levels of the major, naturally occurring cytokinins in Pisum sativum ribulose-1,5-bisphosphate carboxylase small subunit promoter-isopentenyl transferase gene (Pssuipt) -transformed tobacco (Nicotiana tabacum L.) callus were quantified using electrospray-liquid chromatography-tandem mass spectrometry during a 6-week subcultivation period. An ipt gene was expressed under control of a tetracycline-inducible promoter for a more detailed study of cytokinin accumulation and metabolism. Activation of the ipt in both expression systems resulted in the production of mainly zeatin-type cytokinins. No accumulation of isopentenyladenine or isopentenyladenosine was observed. in Pssuipt-transformed calli, as well as in the tetracycline-inducible ipt leaves, metabolic inactivation occurred through O-glucoside conjugation. No significant elevation of cytokinin N-glucosides levels was observed. Side-chain reduction to dihydrozeatin-type cytokinins was observed in both systems. The levels of the endogenous cytokinins varied in time and were subject to homeostatic regulatory mechanisms. Feeding experiments of ipt transgenic callus with (3H) isopentenyladenine and (3H) isopentenyladenosine mainly led to labeled adenine-like compounds, which are degradation products from cytokinin-oxidase activity. Incorporation of radioactivity in zeatin riboside was observed, although to a much lesser extent.

1996

12/3,AB/36 (Item 21 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
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BIOSIS NO.: 199699134083

Chemically induced expression of the rolC-encoded beta-glucosidase in transgenic tobacco plants and analysis of cytokinin metabolism: rolC does not hydrolyze endogenous cytokinin glucosides

AUTHOR: Faiss Martin; Strand Miroslav; Redig Pascale; Dolezal Karel; Hanus Jan; Van Onckelen Harry; Schmuelling Thomas(a)

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JOURNAL: Plant Journal 10 (1):p33-46 **1996**

ISSN: 0960-7412

DOCUMENT TYPE: Article RECORD TYPE: Abstract LANGUAGE: English

ABSTRACT: The rolC gone of Agrobacterium rhizogenes T-DNA plays an essential role in the establishment of hairy root disease and its overexpression in transgenic plants causes pleiotropic developmental alterations. This study investigated whether the biological activity of the rolC beta-glucosidase is due to an alteration of the cytokinin balance in plants. HPLC radiocounting assays of (3H)-labeled cytokinin glucosides fed exogenously to tobacco leaf disks, to rolC expressing Escherichia coli cells or cell-free extracts showed that cytokinin N3- and O-glucosides are the preferred substrate of the rolC protein. Hydrolysis of N7- and N9-glucosides was not detected at substrate concentrations close to physiological levels. Furthermore, these conjugates were also not active as cytokinins in biotests when fed to rolC-expressing tissues. For analysis of the rolC activity on endogenous cytokinin conjugates the gone was expressed under the transcriptional control of a modified tetracycline-inducible 35S promoter. This was done to avoid possible interference with secondary effects or plant homeostatic mechanisms which could mask primary in plants events when transgenes are expressed constitutively. No changes in the endogenous pool of different cytokinin glucosides, as determined by a newly developed electrospray tandem mass spectroscopy directly coupled to high performance liquid chromatography, were found following chemical induction of the rolC gone. Also the levels of free cytokinins remained unchanged after gone induction. Hybrid tobacco plants expressing the cytokinin-synthezising ipt gone and the rolC gone showed added phenotypes indicating that the rolC phenotype is mediated on a signalling pathway different from those of cytokinins . Rolc/ipt hybrids also accumulated high levels of cytokinin O-glucosides. It is concluded that the phenotypic alterations caused by the rolC gone product are not due to a release of free cytokinins from inactive conjugates, most likely because of subcellular compartmentation of the putative substrate.

1996

12/3,AB/37 (Item 22 from file: 5) DIALOG(R)File 5:Biosis Previews(R) (c) 2002 BIOSIS. All rts. reserv.

10424627 BIOSIS NO.: 199699045772

Cytokinin and IAA content in tobacco regenerations carrying the active Agrobacterium ipt gene.

AUTHOR: Makarova R V; Borisova T A; Kefeli V I

AUTHOR ADDRESS: K.A. Timiryazev Inst. Plant Physiol., Russ. Acad. Sci.,

Moscow**Russia

JOURNAL: Doklady Akademii Nauk 346 (2):p280-283 1996

ISSN: 0869-5652

DOCUMENT TYPE: Article RECORD TYPE: Citation

LANGUAGE: Russian; Non-English

1996

12/3,AB/38 (Item 23 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
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10369360 BIOSIS NO.: 199698824278

Analysis of cytokinin biosynthetic gene expression in

transgenic tobacco plants.

AUTHOR: Ma Qing-Hu

AUTHOR ADDRESS: Inst. Bot., Acad. Sinica, Beijing 100044**China

JOURNAL: Chinese Journal of Botany 7 (2):p104-108 1995

ISSN: 1001-0718

DOCUMENT TYPE: Article RECORD TYPE: Abstract LANGUAGE: English

ABSTRACT: The cytokinin biosynthetic gene coding for isopentenyl transferase (ipt) was cloned with its native promoter from Agrobacterium tumefaciens pTiAch58 and introduced into tobacco plants. To overcome the elevated cytokinin levels in suppressing the formation of roots, indolebutyric acid (IBA) was applied to regenerate morphologically normal transgenic tobacco plants. Northern hybridization revealed that the ipt mRNA level in these rooting plants were much lower than those in the primary transformed turnout tissues, and the root was the part in which the ipt gene mRNA level was the lowest in the plant. The determination of endogenous zeatin and zeatin riboside levels gave the same trend with the northern hybridization. These data suggest that the transgenic plants we obtained are a good model for studying the function and regulation of cytokinin in the whole plant

1995

12/3,AB/39 (Item 24 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
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10342649 BIOSIS NO.: 199698797567

Agrobacterium-mediated transformation of apple (Malus x domestica Borkh.):
An assessment of factors affecting regeneration of transgenic
plants.

AUTHOR: De Bondt An; Eggermont Kristel; Penninckx Iris; Goderis Inge; Broekaert Willem F(a)

AUTHOR ADDRESS: (a) F.A. Janssens, Laboratorium voor Genetica, Katholieke Universiteit Leuven, Willem de Croylaan 42**Belgium

JOURNAL: Plant Cell Reports 15 (7):p549-554 1996

ISSN: 0721-7714

DOCUMENT TYPE: Article RECORD TYPE: Abstract LANGUAGE: English

ABSTRACT: We have previously developed a protocol for efficient gene transfer and regeneration of transgenic calli following cocultivation of apple (cv. Jonagold) explants with Agrobacterium tumefaciens (De Bondt et al. 1994, Plant Cell Reports 13: 587-593). Now we report on the optimization of postcultivation conditions for efficient and reproducible regeneration of transgenic shoots from the apple cultivar Jonagold.

Factors which were found to be essential for efficient shoot regeneration were the use of gelrite as a gelling agent and the use of the cytokinin-mimicing thidiazuron in the selective postcultivation medium. Improved transformation efficiencies were obtained by combining the hormones thidiazuron and zeatin and by using leaf explants from in vitro grown shoots not older than 4 weeks after multiplication. Attempts to use phosphinothricin acetyl transferase as a selectable marker were not successful. Using selection on kanamycin under optimal postcultivation conditions, about 2% of the leaf explants developed transgenic shoots or shoot clusters. The presence and expression of the transferred genes was verified by beta-glucuronidase assays and Southern analysis. The transformation procedure has also been successfully applied to several other apple cultivars.

1996

12/3,AB/40 (Item 25 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
(c) 2002 BIOSIS. All rts. reserv.

10289005 BIOSIS NO.: 199698743923
Effect of heat shock on the dynamics of cytokinin concentration in transgenic and intact tobacco plants.

AUTHOR: Veselov S Yu; Kudoyarova G R(a); Mustafina A R; Valcke R AUTHOR ADDRESS: (a) Inst. Biol., Ufa Res. Cent., Russ. Acad. Sci., prospekt Oktyabrya 69, 450054 Ufa**Russia

JOURNAL: Fiziologiya Rastenii (Moscow) 42 (5):p696-699 1995

ISSN: 0015-3303

DOCUMENT TYPE: Article RECORD TYPE: Abstract

LANGUAGE: Russian; Non-English

SUMMARY LANGUAGE: Russian; Non-English

ABSTRACT: The daily dynamics of the concentration of endogenous cytokinins was determined in transgenic Nicotiana tabacum shoots. In these plants, the expression of ipt-gene controlling isopentenyltransferase synthesis was induced by heat shock. Incubation of transgenic plants during one hour at 40 degree C resulted in an increase in endogenous cytokinin concentration as compared to the control transgenic plants constantly exposed to the temperature of 24 degree C. However, this increase lasted only for a short period of time and no differences were observed between the experimental and control plants 5 hours after the effect of heat shock. In the shoots of intact plant seedlings, heat shock induced the activation of processes focused at a decrease in cytokinin concentration. This phenomenon can be considered a natural reaction of plants to heat-induced stress. The realization of this natural reaction in transgenic plants can be a reason for a short duration of cytokinin accumulation induced by heat shock.

1995

12/3,AB/41 (Item 26 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
(c) 2002 BIOSIS. All rts. reserv.

10199993 BIOSIS NO.: 199698654911

Cytokinin oxidase-purification by affinity chromatography and

activation by caffeic acid. AUTHOR: Wang J; Letham D S(a)

AUTHOR ADDRESS: (a) Cooperative Res. Cent. Plant Sci., Res. Sch. Biol. Sci., Australian Natl. Univ., P.O. Box 475, C**Australia

JOURNAL: Plant Science (Shannon) 112 (2):p161-166 1995

ISSN: 0168-9452

DOCUMENT TYPE: Article RECORD TYPE: Abstract LANGUAGE: English

ABSTRACT: Two cytokinin analogues, 6-di-(isopent-2-enyl)aminopurine and 6-(N-isopent-2-enyl-N-methylamino)purine were found to be effective inhibitors of cytokinin oxidase prepared from tobacco tissue cultures expressing the cytokinin biosynthesis gene ipt. The latter analogue was conjugated at the N-9 position to Sepharose through a 12-atom spacer moiety. This yielded a matrix for preparation of an affinity column for further purification of cytokinin oxidase that had been partially purified by other methods. The activity of cytokinin oxidase was enhanced by caffeic acid and to a lesser extent by other phenolic compounds tested.

1995

12/3,AB/42 (Item 27 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
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10157530 BIOSIS NO.: 199698612448

The effect of an elevated **cytokinin** level using the **ipt** gene and N-6-benzyladenine on single node and intact **plant** tuberization

AUTHOR: Galis Vvan Jiri Macas(a); Vlasak Josef; Ondrej Milos; Van Onckelen Henri A

AUTHOR ADDRESS: (a) Inst. Plant Molecular Biol., Acad. Sci., Czech Republic, Branisovska 31, 370 05 Ceske Budejovice**Czech Republic

JOURNAL: Journal of Plant Growth Regulation 14 (3):p143-150 1995

ISSN: 0721-7595

DOCUMENT TYPE: Article RECORD TYPE: Abstract LANGUAGE: English

ABSTRACT: Two models of potato (Solanum tuberosum L.) tuberization in vitro (intact plants and single nodes) were used to study the role of cytokinins in this process. We applied hormone in two different ways. The exogenous addition of 10 mg cntdot L-1 N-6-benzyladenine (BA) into the tuberization medium resulted in advanced tuber formation in intact plants, and microtubers appeared 10-20 days earlier than in the experiments in which no cytokinin was supplied. Transformation with the Agrobacterium tumefaciens ipt gene provided potato clones with endogenously elevated cytokinin levels (3-20 times higher zeatin riboside content in different clones). The onset of tuberization in intact ipt-transformed plants with low transgene expression was advanced in comparison with control material, and exogenously applied BA further promoted the tuberization process. On the contrary, tuberization was strongly inhibited in ipt -transformed nodes, and an external increase of the cytokinin level caused complete inhibition of explant growth. In untransformed (control) nodes cytokinin application resulted in primary and secondary tuber formation, which depended on the BA concentration in cultivation media.

1995

12/3,AB/43 (Item 28 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
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10003903 BIOSIS NO.: 199598458821

Expression of a wound-inducible cytokinin biosynthesis gene in transgenic tobacco: Correlation of root expression with induction of cytokinin effects.

AUTHOR: Smigocki Ann C

AUTHOR ADDRESS: Plant Molecular Biology Lab., USDA/ARS, 10300 Baltimore

Ave., Build. 006, Room 118, Beltsville, MD 2**USA

JOURNAL: Plant Science (Limerick) 109 (2):p153-163 1995

ISSN: 0168-9452

DOCUMENT TYPE: Article RECORD TYPE: Abstract LANGUAGE: English

ABSTRACT: The Agrobacterium-derived cytokinin-biosynthesis gene ipt was fused to the wound-inducible proteinase-inhibitor-IIK gene promoter from potato and introduced into Nicotiana plumbaginifolia and N. tabacum. Maximum accumulation of ipt transcripts in the leaves of transgenic plants was observed within 3-24 h after leaf wounding. Root and stem ipt messages were not detected in unwounded transgenic N. plumbaginifolia PI-II-ipt seedlings until after the plants bolted whereas in N. tabacum, a relatively low level of root and stem expression was evident only prior to stem elongation and not detected after the plants bolted. Atypical cytokinin effects were observed with the N. plumbaginifolia but not N. tabacum transformants. Transgenic N. plumbaginifolia plants bolted sooner, were taller than control plants and had larger leaves with lower specific fresh weights and chlorophyll content. At flowering, the emergence of numerous lateral shoots from lower stem sections and basal leaf greening followed the moderate increase in root ipt transcripts and corresponded to a greater than 100-fold increase in zeatin and zeatinriboside cytokinin concentrations. The expression pattern of the PI-II-ipt gene followed that of the PI-IIK gene and, when expressed in the root, corresponded with induction of characteristic cytokinin effects.

1995

12/3,AB/44 (Item 29 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
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09975304 BIOSIS NO.: 199598430222

Effect of Auxin on Expression of the Isopentenyl

Transferase Gene (ipt) in Transformed Bean (Phaseolus

vulgaris L.) Single-Cell Clones Induced by Agrobacterium tumefaciens C58. AUTHOR: Song Jai Young; Choi Eun Yeung; Lee Hyeun Se; Choi Dong-Woog; Oh

Man-Ho; Kim Sang-Gu(a)

AUTHOR ADDRESS: (a)Dep. Biol. Res. Cent. Cell Differentiation, Seoul Natl. Univ., Seoul 151-742**South Korea

JOURNAL: Journal of Plant Physiology 146 (1-2):p148-154 1995 ISSN: 0176-1617

DOCUMENT TYPE: Article RECORD TYPE: Abstract LANGUAGE: English

ABSTRACT: The effect of auxin on the endogenous cytokinin content and on the expression of isopentenyl transferase gene (ipt) was investigated in bean (Phaseolus vulgaris L. cv. Palgong) tumor single-cell clones induced by Agrobacterium tumefaciens C58. The major endogenous cytokinins of tumor single-cell clones were zeatin and zeatin riboside. Endogenous zeatin and zeatin riboside levels in tumor single-cell clones cultured on an auxin-supplemented medium were

reduced by six-fold and eight-fold, respectively, while tumor single-cell clones cultured on the 5.0 mu-M kinetin-supplemented medium did not exhibit any reduction in the levels of these cytokinins. The mRNAs isolated from normal single-cell clones cultured on 5.0 mu-M kinetin and 2.5 mu-M picloram-supplemented medium, from transformed single-cell clones cultured on hormone-free medium, and from transformed single-cell clones cultured on 2.5 mu-M picloram-supplemented medium, were subjected to Northern blot hybridization. The ipt transcript was not detected in tumor single-cell clones cultured on picloram-supplemented medium, but the ipt mRNA was detected in tumor single-cell clones cultured on hormone-free medium. The amount of ipt mRNA in tumor single-cell clones was found to decrease with time in cultures grown on picloram-supplemented medium. The nopaline synthase gene (nos) transcript was detected in the tumor single-cell clones from both culture conditions. It is concluded that picloram regulates the ipt mRNA steady state level, either at the transcriptional level or by affecting ipt mRNA stability.

1995

12/3,AB/45 (Item 30 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
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09832302 BIOSIS NO.: 199598287220

The effect of auxin on cytokinin levels and metabolism in transgenic tobacco tissue expressing an ipt gene.

AUTHOR: Zhang R; Zhang X; Wang J; Letham D S(a); McKinney S A; Higgins T J

AUTHOR ADDRESS: (a) Cooperative Res. Cent. Plant Sci., Australian Natl.

Univ., PO Box 475, Canberra, ACT 2601**Australia JOURNAL: Planta (Heidelberg) 196 (1):p84-94 1995

ISSN: 0032-0935

DOCUMENT TYPE: Article RECORD TYPE: Abstract LANGUAGE: English

ABSTRACT: The ipt gene from the T-DNA of Agrobacterium tumefaciens was transferred to tobacco (Nicotiana tabacum L.) in order to study the control which auxin appears to exert over levels of cytokinin generated by expression of this gene. The transgenic tissues contained elevated levels of cytokinins, exhibited cytokinin and auxin autonomy and grew as shooty calli on hormone-free media. Addition of 1-naphthylacetic acid to this culture medium reduced the total level of cytokinins by 84% while 6-benzylaminopurine elevated the cytokinin level when added to media containing auxin. The cytokinins in the transgenic tissue were labelled with 3H and auxin was found to promote conversion of zeatin-type cytokinins to 3H-labelled adenine derivatives. When the very rapid metabolism of exogenous (3H) zeatin riboside was suppressed by a phenylurea derivative, a noncompetitive inhibitor of cytokinin oxidase, auxin promoted metabolism to adenine-type compounds. Since these results indicated that auxin promoted cytokinin oxidase activity in the transformed tissue, this enzyme was purified from the tobacco tissue cultures. Auxin did not increase the level of the enzyme per unit tissue protein, but did enhance the activity of the enzyme in vitro and promoted the activity of both glycosylated and non-glycosylated forms. This enhancement could contribute to the decrease in cytokinin level induced by auxin. Studies of cytokinin biosynthesis in the transgenic tissues indicated that transhydroxylation of isopentenyladenine-type cytokinins to yield zeatin-type cytokinins occurred principally at the nucleotide level.

12/3,AB/46 (Item 31 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
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09765648 BIOSIS NO.: 199598220566

Increase of endogenous zeatin riboside by introduction of the ipt gene in wild type and the lateral suppressor mutant of tomato.

AUTHOR: Groot Steven P C(a); Bouwer Reinoud(a); Busscher Marco(a); Lindhout Pim; Dons Hans J(a)

AUTHOR ADDRESS: (a) Dep. Dev. Biol., Cent. Plant Breed. Reprod. Res., P.O. Box 16, NL-6700 AA Wageningen**Netherlands

JOURNAL: Plant Growth Regulation 16 (1):p27-36 1995

ISSN: 0167-6903

DOCUMENT TYPE: Article RECORD TYPE: Abstract LANGUAGE: English

ABSTRACT: We studied axillary meristem formation of the lateral suppressor (ls) mutant of tomato after elevating the endogenous cytokinin levels through introduction of the isopentenyltransferase (ipt) gene from Agrobacterium tumefaciens. Growth and development of several transformants were examined during in vitro culture. Transformants exhibited phenotypes varying in severity and were divided into four classes. A number of the ipt transformants had a normal phenotype, as non-transformed plants. Others showed a mild to severe ' cytokinin-like' phenotype. Transformants with a mild phenotype exhibited reduced internode length and reduced root development. Transformants with a severe phenotype showed even shorter internodes, loss of apical dominance, reduction of leaf size, production of callus at the basis of the shoots and absence of root development or development of green non-branching roots. The severity of the phenotype correlated well with the level of ipt gene expression, as measured by northern analysis. Transformants with a severe phenotype also exhibited increased levels of zeatin riboside, but zeatin levels were not elevated. The increase in endogenous zeatin riboside levels in the 1s mutant did not restore axillary meristem formation, but sometimes bulbous structures were formed in the initially 'empty' leaf axils. Several adventitious meristems and shoots developed from below the surface of these structures. It is concluded that a reduced level of cytokinins in the 1s mutant shoots is not responsible for the absence of axillary meristem formation.

1995

12/3,AB/47 (Item 32 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
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09716852 BIOSIS NO.: 199598171770

Production of high solids tomatoes through molecular modification of levels of the plant growth regulator cytokinin.

AUTHOR: Martineau Belinda(a); Summerfelt Kristin R; Adams Dawn F; Deverna Joseeph W

AUTHOR ADDRESS: (a) Calgene Inc., 1920 Fifth St., Davis, CA 95616**USA

JOURNAL: Bio-Technology (New York) 13 (3):p250-254 1995

ISSN: 0733-222X

DOCUMENT TYPE: Article RECORD TYPE: Abstract LANGUAGE: English ABSTRACT: Chimeric isopentenyl transferase (ipt) gene constructs were prepared and introduced into tomato plants via Agrobacterium-mediated transformation. Expression of the ipt gene, which encodes a key enzyme involved in the biosynthesis of the plant growth regulator cytokinin, was modulated using a promoter from a gene expressed primarily in tomato ovaries. As expected, the ipt gene was expressed, and levels of cytokinin were increased, in ovaries of the transgenic plants. Plant yield and fruit processing characteristics of these transgenic plants were examined during three consecutive years of field testing. Levels of total solids were significantly increased in six of seven, and soluble solids were significantly increased in five of seven, independent transgenic tomato lines.

1995

12/3,AB/48 (Item 33 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
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09536881 BIOSIS NO.: 199497545251

Specific expression of isopentenyl transferase gene in

transgenic tobacco.

AUTHOR: Ma Qin-Hu(a); Zhang; Ren; Higgins Thomas J V

AUTHOR ADDRESS: (a) Inst. Botany, Acadmia Sinica, Beijing 100044**China

JOURNAL: Acta Botanica Sinica 36 (5):p339-344 1994

ISSN: 0577-7496

DOCUMENT TYPE: Article RECORD TYPE: Abstract

LANGUAGE: Chinese; Non-English SUMMARY LANGUAGE: Chinese; English

ABSTRACT: The cytokinin biosynthetic gene, isopentenyl transferase (ipt) gene of Agrobacterium tumefaciens was fused to a petunia ribulose bisphosphate carboxylase small subunit (SSU301) promoter and introduced into tobacco plants. The expression pattern of this chimeric SSU-ipt gene was studied in the transgenic plants, and the endogenous levels of cytokinins were determined. It was revealed that ipt mRNA level was increased in light-cultured transgenic tobacco tissues, but was undetectable in dark cultured condition. The levels of zeatin and zeatin riboside in the transgenic tissues in light increased about 10 times as compared with the tissues in dark. The results show that the petunia SSU301 promoter can specifically direct the expression of the ipt gene in the transgenic tobacco. These SSU-ipt gene transgenic tobacco plants will provide valuable materials for studies of cytokinin's functions in phytosynthetic tissues and in the light-related physiological processes.

1994

12/3,AB/49 (Item 34 from file: 5) DIALOG(R)File 5:Biosis Previews(R) (c) 2002 BIOSIS. All rts. reserv.

09446154 BIOSIS NO.: 199497454524

Transgenic tobacco plants that overproduce cytokinins show increased tolerance to exogenous auxin and auxin transport inhibitors.

AUTHOR: Li Yi; Shi Xiangyang; Strabala Timothy J; Hagen Gretchen; Guilfoyle Tom J(a)

AUTHOR ADDRESS: (a) Dep. Biochem., 117 Schweitzer Hall, Univ. Missouri,

Columbia, MO 65211**USA

JOURNAL: Plant Science (Limerick) 100 (1):p9-14 1994

ISSN: 0168-9452

DOCUMENT TYPE: Article RECORD TYPE: Abstract LANGUAGE: English

ABSTRACT: Transgenic tobacco plants expressing the Agrobacterium tumefaciens cytokinin biosynthetic ipt gene under the control of an auxin-inducible SAUR (Small Auxin-Up RNA) gene promoter were used to study interactions between exogenously applied auxins or auxin transport inhibitors and endogenously produced cytokinins. The transgenic plants used in this study had cytokinin levels about 10-fold higher than non-transformed tobacco plants. In aseptic culture, the transgenic tobacco plants exhibited increased tolerance to the toxic effects of high concentrations of exogenously applied auxins. This tolerance is exemplified by increased plant height and fresh weight in transgenic plants treated with auxin compared to similarly treated non-transformed plants. In addition to increased tolerance to exogenous auxins, the transgenic plants showed increased tolerance to applied auxin transport inhibitors.

1994

12/3,AB/50 (Item 35 from file: 5) DIALOG(R)File 5:Biosis Previews(R) (c) 2002 BIOSIS. All rts. reserv.

09380286 BIOSIS NO.: 199497388656 Phenotype modification and disease resistance via regulated expression of the cytokinin biosynthesis gene.

AUTHOR: Smigocki Ann C

AUTHOR ADDRESS: Plant Mol. Biol. Lab., ARS/USDA, Beltsville, MD 20705-2350 **USA

JOURNAL: Hortscience 29 (5):p574 1994

CONFERENCE/MEETING: 91st Annual Meeting of the American Society for Horticultural Science Corvallis, Oregon, USA August 7-10, 1994

ISSN: 0018-5345

RECORD TYPE: Citation LANGUAGE: English

1994

12/3,AB/51 (Item 36 from file: 5) DIALOG(R)File 5:Biosis Previews (R) (c) 2002 BIOSIS. All rts. reserv.

09344903 BIOSIS NO.: 199497353273

Osmotic stress symptoms in transgenic tobacco expressing ipt

from A. tumefaciens.

AUTHOR: Thomas John C(a); Smigocki Ann C; Bohnert Hans J AUTHOR ADDRESS: (a)Dep. Biochem., Univ. Arizona, Tucson, AZ 85721**USA JOURNAL: Plant Physiology (Rockville) 105 (1 SUPPL.):p71 1994 CONFERENCE/MEETING: Annual Meeting of the American Society of Plant Physiologists Portland, Oregon, USA July 30-August 3, 1994

ISSN: 0032-0889

RECORD TYPE: Citation LANGUAGE: English

1994

DIALOG(R) File 5:Biosis Previews (R) (c) 2002 BIOSIS. All rts. reserv.

09261434 BIOSIS NO.: 199497269804

Fruit-specific expression of the A. tumefaciens isopentenyl transferase gene in tomato: Effects on fruit ripening and defense-related gene expression in leaves.

AUTHOR: Martineau Belinda(a); Houck Catherine M; Sheehy Raymond E; Hiatt

AUTHOR ADDRESS: (a) Calgene Fresh Inc., 1910 Fifth St., David, CA 95616**USA

JOURNAL: Plant Journal 5 (1):p11-19 1994

ISSN: 0960-7412

DOCUMENT TYPE: Article RECORD TYPE: Abstract LANGUAGE: English

ABSTRACT: This paper describes the analysis of tomato plants transformed with a chimeric gene consisting of the promoter region of a fruit specifically expressed tomato gene linked to the ipt gene coding sequences from the Ti plasmid of Agrobacterium tumefaciens. The pattern of expression of this chimeric gene was found to be consistent with the expression of the endogenous fruit-specific gene and consequently, plants expressing the chimeric gene were phenotypically normal until fruit maturation and ripening. A dramatically altered fruit phenotype, islands of green pericarp tissue remaining on otherwise deep red ripe fruit, was then evident in many of the transformed plants. Cytokinin levels in transformed plant fruit tissues were 10 to 100-fold higher than in control fruit. In the leaves of a fruitbearing transformant, despite a lack of detectable ipt mRNA accumulation, approximately fourfold higher than control leaf levels of cytokinin were detected. It is suggested that cytokinin produced in fruit is being transported to the leaves since accumulation in leaves of PR-1 and chitinase mRNAs, which encode defense-related proteins known to be induced by cytokinin, occurred only when the transformant was reproductively active. Effects of elevated cytokinin levels on tomato fruit gene expression and cellular differentiation processes are also described.

1994

12/3,AB/53 (Item 38 from file: 5) DIALOG(R)File 5:Biosis Previews(R) (c) 2002 BIOSIS. All rts. reserv.

BIOSIS NO.: 199497197215

Cytokinins modulate stress response genes in isopentenyl transferase-transformed Nicotiana plumbaginifolia plants.

AUTHOR: Harding S A; Smigocki A C(a)

AUTHOR ADDRESS: (a) Plant Mol. Biol. Lab., USDA/ARS Beltsville, MD 20705**

JOURNAL: Physiologia Plantarum 90 (2):p327-333 1994

ISSN: 0031-9317

DOCUMENT TYPE: Article RECORD TYPE: Abstract LANGUAGE: English

ABSTRACT: The effects of transiently elevated cytokinin levels on gene expression following stress were examined in transgenic Nicotiana plumbaginifolia plants. Plants were transformed with a bacterial gene encoding isopentenyl transferase (ipt) cloned behind the heat shock (HS) protein 70 promoter from Drosophila melanogaster. Following a 1-h, 45 degree C HS of whole plants, the ipt transcript levels in leaves increased 30- to 40-fold. Analysis of in vitro translation products of leaf messenger RNA showed rapid isopentenyl transferase-dependent changes in gene expression. A subset comprising 1 to 2% of resolvable translation products was specifically up-regulated in heat shock ipt-inducible (HS-ipt) plants. Several cDNA clones were isolated which correspond to mRNAs that are up-regulated 2- to 4-fold in HS-ipt plants. Two of the cDNAs encode stress-related polypeptides, one a member of a class of small heat shock polypeptides (HSP) and the other, a wound-inducible glycine-rich protein. Benzylaminopurine feeding experiments show that the HSP transcripts are up-regulated by other treatments including watering but that cytokinins strongly accelerate or amplify the response. These data are the first to show altered modulation of stress-induced genes in intact plants transformed with the cytokinin biosynthesis gene ipt.

1994

12/3,AB/54 (Item 39 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
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09138338 BIOSIS NO.: 199497146708

The role of hormones in apical dominance. New approaches to an old problem in **plant** development.

AUTHOR: Cline Morris G

AUTHOR ADDRESS: Dep. Plant Biol., Ohio State Univ., 1735 Neil Ave.,

Columbus, OH 43210**USA

JOURNAL: Physiologia Plantarum 90 (1):p230-237 1994

ISSN: 0031-9317

DOCUMENT TYPE: Literature Review

RECORD TYPE: Abstract LANGUAGE: English

ABSTRACT: The role of hormones in apical dominance has been under investigation with traditional 'spray and weigh' methods for nearly 5 decades. Even though the precision of hormone content analyses in tissue has greatly improved in recent years, there have been no significant breakthroughs in our understanding of the action mechanism of this classical developmental response. Auxin appears to inhibit axillary bud outgrowth whereas cytokinins will often promote it. Conclusive evidence for a direct role of these or other hormones in apical dominance has not been forthcoming. However, promising new tools and approaches recently have begun to be utilized. The manipulation of endogenous hormone levels via the use of transgenic plants transformed with bacterial genes (iaaM and ipt from Agrobacterium tumefaciens and iaaL from Pseudomonas syringae pv. savastanoi) has demonstrated powerful effects of auxin and cytokinin on axillary bud outgrowth. Also, possible auxin and cytokinin involvement of rolB and C genes from Agrobacterium rhizogenes whose activity is associated with reduced apical dominance in dicotyledons has received considerable attention. The characterization of unique mRNAs and proteins in non-growing and growing lateral buds before and after apical dominance release is helping to lay the groundwork for the elucidation of signal transduction and cell cycle regulation in this response. The use of auxin-deficient, and auxin/ethylene-resistant mutants has provided another approach for analyzing the role of these hormones. The presumed eventual employment of molecular assay systems (SAUR/GH3 promoters fused with GUS reporter gene) which are presently being developed for analyzing auxin localized in lateral buds will hopefully provide a critical test for the direct auxin inhibition hypothesis.

12/3,AB/55 (Item 40 from file: 5) DIALOG(R)File 5:Biosis Previews(R) (c) 2002 BIOSIS. All rts. reserv.

BIOSIS NO.: 199497089575

Morphometric analysis of the growth of Phsp70-ipt transgenic tobacco plants.

AUTHOR: Van Loven Karen; Beinsberger Susy E I; Valcke Roland L M(a); Van Onckelen Henri A; Clijsters Herman M M

AUTHOR ADDRESS: (a) Limburgs Universitair Centrum, Dept. SBG, Universitaire Campus, B-3590 Diepenbeek**Belgium

JOURNAL: Journal of Experimental Botany 44 (268):p1671-1678 1993

ISSN: 0022-0957

DOCUMENT TYPE: Article RECORD TYPE: Abstract LANGUAGE: English

ABSTRACT: The effect of introducing a supplementary ipt-gene into the genome of Nicotiana tabacum L. cv. Petit Havana SR1 is studied on the morphological plant development. The ipt-gene, accounting for the biosynthesis of cytokinins, was coupled to the heat-inducible hsp70- promoter from Drosophila melanogaster. Besides the influence of the hormonal changes involved, the effects of the experimental conditions are examined, namely the in vitro growth conditions for selecting transformed plants and the heat treatment to induce ipt-gene expression. The phenotype of the plants is determined by the tissue sensitivity to three factors: (1) heat treatment reduces stem elongation and diameter growth; (2) in vitro pre-cultivation also reduces stem elongation; and (3) expression of the ipt-gene stimulates diameter growth, induces debudding of the axillary shoots and inhibits root development. In addition, axillary bud development indicates that in vitro cultivation affects ipt-gene expression.

1993

12/3,AB/56 (Item 41 from file: 5) DIALOG(R)File 5:Biosis Previews(R) (c) 2002 BIOSIS. All rts. reserv.

09042189 BIOSIS NO.: 199497050559

Alterations in auxin and cytokinin metabolism of higher plants due to expression of specific genes from pathogenic bacteria: A review.

AUTHOR: Hamill John D

AUTHOR ADDRESS: Dep. Genet. Dev. Biol., Monash Univ., Clayton, VIC 3168** Australia

JOURNAL: Australian Journal of Plant Physiology 20 (4-5):p405-423 1993

ISSN: 0310-7841

DOCUMENT TYPE: Literature Review

RECORD TYPE: Abstract LANGUAGE: English

ABSTRACT: This review deals with the physiological and morphological effects of altering the auxin/cytokinin balance in transgenic plants by expressing specific genes from pathogenic bacteria. Genes which have been used to alter auxin levels or sensitivity in transgenic **plants** include the iaaM/iaaH genes from Agrobacterium tumefaciens and A. rhizogenes; gene 5 and possibly gene 6b from A.

tumefaciens; the rol B and possibly the rol A gene from A. rhizogenes and the iaaL gene from Pseudomonas syringae subsp. savastanoi (P. savastanoi). Genes which have been used to alter cytokinin levels in transgenic plants include the ipt gene from A. tumefaciens and the rol C gene from A. rhizogenes. A variety of biochemical mechanisms have been identified which result in alterations to phytohormone levels following expression of these genes in transgenic plants. Many of the effects on plant development are consistent with observations made following exogenous auxin and/or cytokinin application to plant tissues, and the availability of these genes offers a new approach to the study of plant physiology using transformation methodology.

1993

12/3,AB/57 (Item 42 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
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08936028 BIOSIS NO.: 199396087529

Expression of a cytokinin synthesis gene from Agrobacterium tumefaciens T-DNA in basket willow (Salix viminalis).

AUTHOR: Vahala T(a); Eriksson T; Tillberg E; Nicander B

AUTHOR ADDRESS: (a) Dep. Molecular Genetics, Swedish Univ. Agricultural

Sciences, Box 7003, S-75006 Uppsala**Sweden

JOURNAL: Physiologia Plantarum 88 (3):p439-445 1993

ISSN: 0031-9317

DOCUMENT TYPE: Article RECORD TYPE: Abstract LANGUAGE: English

ABSTRACT: Willow cells transformed with an **ipt** gene from Agrobacterium tumefaciens grow in tissue culture as undifferentiated callus without shoot induction. We show that the transformed calluses contained high levels of the **cytokinins** 9-beta-D-ribofuranosyl zeatin and its monophosphate, demonstrating the presence of a functional **isopentenyl transferase** enzyme. The **ipt** gene was transcribed at different levels in different transformed callus fines. The absence of shoot differentiation is apparently not due to a lack of zeatin-type **cytokinins** in the transformed callus.

1993

12/3,AB/58 (Item 43 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
(c) 2002 BIOSIS. All rts. reserv.

08514826 BIOSIS NO.: 199344064826 Cloning and expression of the IPT gene.

AUTHOR: Barbour Sandra L; Schaff D A; Frett J J

AUTHOR ADDRESS: Dep. Plant Soil Sci., Univ. Delaware, Newark, DE 19717**

JOURNAL: Hortscience 27 (11):p1160 1992

CONFERENCE/MEETING: Annual Meeting of the ASHS (American Society for Horticultural Science) Northeast Region Amherst, Massachusetts, USA January 9-11, 1992

ISSN: 0018-5345 RECORD TYPE: Citation LANGUAGE: English

1992

12/3,AB/59 (Item 44 from file: 5) DIALOG(R) File 5:Biosis Previews(R) (c) 2002 BIOSIS. All rts. reserv.

08140288 BIOSIS NO.: 000093127436 FASCIATION INDUCTION BY THE PHYTOPATHOGEN RHODOCOCCUS-FASCIANS DEPENDS UPON UPON A LINEAR PLASMID ENCODING A CYTOKININ SYNTHASE GENE AUTHOR: CRESPI M; MESSENS E; CAPLAN A B; VAN MONTAGU M; DESOMER J AUTHOR ADDRESS: LABORATORIUM VOOR GENETICA, UNIVERSITEIT GENT, B-9000 GENT, BELGIUM.

JOURNAL: EMBO (EUR MOL BIOL ORGAN) J 11 (3). 1992. 795-804. 1992 FULL JOURNAL NAME: EMBO (European Molecular Biology Organization) Journal CODEN: EMJOD

RECORD TYPE: Abstract LANGUAGE: ENGLISH

ABSTRACT: Rhodococcus fascians is a nocardiform bacteria that induces leafy galls (fasciation) of dicotyledonous and several monocotyledonous plants. The wild-type strain D188 contained a conjugative, 200 kb linear extrachromosomal element, pFiD188. Linear plasmid-cured strains were avirulent and reintroduction of this linear element restored virulence. Pulsed field electrophoresis indicated that the chromosome might also be a linear molecule of 4 megabases. Three loci involved in phytopathogenicity have been identified by insertion mutagenesis of this Fi plasmid. Inactivation of the fas locus resulted in avirulent strains, whereas insertions in the two other loci affected the degree of virulence, yielding attenuated (att) and hypervirulent (hyp) bacteria. One of the genes within the fas locus encoded an isopentenyltransferase (IPT) with low homology to analogous proteins from Gram-negative phytopathogenic bacteria. IPT activity was detected after expression of this protein in Escherichia coli cells. In R. fascians, ipt expression could only be detected in bacteria induced with extracts from fasciated tissue. R. fascians strains without the linear plasmid but containing this fas locus alone could not provoke any phenotype on plants, indicating additional genes from the linear plasmid were also essential for virulence. These studies, the first genetic analysis of the interaction of a Gram-positive bacterium with plants, suggest that a novel mechanism for plant tumour induction has evolved in R. fascians independently from the other branches of the eubacteria.

1992

12/3,AB/60 (Item 45 from file: 5) DIALOG(R)File 5:Biosis Previews(R)

(c) 2002 BIOSIS. All rts. reserv.

07949577 BIOSIS NO.: 000093028675 REGENERATION OF PLANTS FROM PEACH EMBRYO CELLS INFECTED WITH A SHOOTY MUTANT STRAIN OF AGROBACTERIUM

AUTHOR: SMIGOCKI A C; HAMMERSCHLAG F A

AUTHOR ADDRESS: U.S. DEP. AGRIC., AGRIC. RES. SERV., PLANT MOL. BIOL. LAB., BELTSVILLE, MD. 20705, USA.

JOURNAL: J AM SOC HORTIC SCI 116 (6). 1991. 1092-1097. 1991

FULL JOURNAL NAME: Journal of the American Society for Horticultural

Science

CODEN: JOSHB

RECORD TYPE: Abstract

LANGUAGE: ENGLISH

ABSTRACT: Immature 'Redhaven' peach [Prunus persica (L.) Batsch] embryos were infected with a shooty mutant strain of Agrobacterium tumefaciens, tms328::Tn5, which carries an octopine-type Ti plasmid with a functional

cytokinin gene and a mutated auxin gene. Shoots were regenerated from embryo-derived callus that was initiated on MS medium lacking phytohormones. Shoots exhibit increased frequency of branching and were more difficult to root than the noninfected. Transcripts of the tms328::Tn5-cytokinin gene were detected using northern analyses of total plant RNA. Polymerase chain reaction of genomic DNA and cDNA resulted in amplification of DNA fragments specific for the cytokinin gene, as determined by restriction enzyme and Southern analyses. The concentrations of the cytokinins zeatin and zeatin riboside in the leaves of regenerated plants were on the average 51-fold higher than in leaves taken from nontransformed plants. None of the shoots or callus tissues were positive for octopine. The expression of the T-DNA encoded cytokinin gene promotes growth of peach cells in the absence of phytohormones, thus serving as a marker for transformation. In addition, this gene appears to promote morphogenesis without an auxin inductive step.

1991

12/3,AB/61 (Item 46 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
(c) 2002 BIOSIS. All rts. reserv.

07400418 BIOSIS NO.: 000091016028
RESTORATION OF SHOOTY MORPHOLOGY OF A NONTUMOROUS MUTANT OF
NICOTIANA-GLAUCA X NICOTIANA-LANGSDORFFII BY CYTOKININ AND THE
ISOPENTENYLTRANSFERASE GENE

AUTHOR: FENG X-H; DUBE S K; BOTTINO P J; KUNG S-D AUTHOR ADDRESS: CENT. AGRIC. BIOTECHNOL., UNIV. MD., COLLEGE PARK, MD. 20742.

JOURNAL: PLANT MOL BIOL 15 (3). 1990. 407-420. 1990

FULL JOURNAL NAME: Plant Molecular Biology

CODEN: PMBID

RECORD TYPE: Abstract LANGUAGE: ENGLISH

ABSTRACT: The shooty morphology of a nontumorous amphidiploid mutant of Nicotiana glauca Grah. .times. N. langsdorffii Weinm. was restored by cytokinins, whether exogenously applied or endogenously produced by transformation of the mutant with a transfer DNA (T-DNA) cytokinin -biosynthesis gene (isopentenyltransferase; ipt). Auxins alone did not confer this effect. Similar transformation was not achieved for the parental species. In the case of transformation with the ipt gene, selection of the transformed tissues was based on its hormone-independent growth in the presence of the antibiotic kanamycin. Transformed tissues exhibited a shooty morphology, indistinguishable from the of wildtype genetic tumors N. glauca .times. N. langsdorffii. This altered phenotype was caused by the presence and constitutive expression of the ipt gene. the insertion and expression of this gene in transformed tissues was confirmed by using the polymerase chain reaction (PCR) technique as well as conventional molecular hybridization analysis. Expression of the ipt gene led to an elevated level of cytokinin in the transformed mutant tissues. This evidence supports the notion that genetic tumors are caused, at least in part, by elevated levels of cytokinin interspecific hybrids.

1990

12/3,AB/62 (Item 47 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
(c) 2002 BIOSIS. All rts. reserv.

BIOSIS NO.: 000090062686 07282799

AGROBACTERIUM-MEDIATED TRANSFORMATION OF THE CULTIVATED STRAWBERRY FRAGARIA-ANANASSA DUCH. USING DISARMED BINARY VECTORS

AUTHOR: JAMES D J; PASSEY A J; BARBARA D J

AUTHOR ADDRESS: INST. HORTICULTURAL RESEARCH, EAST MALLING, MAIDSTONE, KENT, ME19 6BJ, UK.

JOURNAL: PLANT SCI (LIMERICK) 69 (1). 1990. 79-94. 1990

CODEN: PLSCE

RECORD TYPE: Abstract LANGUAGE: ENGLISH

ABSTRACT: Two disarmed Ti-binary vectors in Agrobacterium tumefaciens have been used to produce viable transgenic strawberry plants. Fertile strawberry plants with a normal phenotype were regenerated after transformation with pBIN6, which carries genes for nopaline synthase (nos) and neomycin phosphotransferase (nptII) (conferring kanamycin resistance). The transfer and expression of the two genes was confirmed by Southern blot analysis, the detection of nopaline synthase (NOS) activity in vegetative and reproductive tissues and rooting in vitro in the presence of kanamycin. The nos gene continued to be expressed in glasshouse-grown plants many months after removal from in vitro growth conditions. After selfing the RO plants nos segregated in the R1 progeny according to a 3:1 Mendelian ratio. In in vitro germinated seedlings there was complete correlation between the presence of nopaline synthase activity and the ability of leaf segments to produce callus on a medium containing kanamycin. Transgenic clones that exhibited an abnormal phenotype associated with cytokinin overproduction were produced when plants were transformed with pSS1, a derivative of pBIN19 carrying both the nptII gene and the ipt gene (encoding the enzyme isopentenyltransferase). Shoots of these clones grew on hormone-free medium, could not be induced to root and their growth was unaffected by the presence of 50 .mu.g/ml kanamycin in hormone-free media.

1990

12/3,AB/63 (Item 48 from file: 5) DIALOG(R)File 5:Biosis Previews(R) (c) 2002 BIOSIS. All rts. reserv.

06925388 BIOSIS NO.: 000089058780

TRANSFER OF THE AGROBACTERIAL CYTOKININ BIOSYNTHESIS GENE INTO TOBACCO PLANTS

AUTHOR: YUSIBOV V M; POGOSYAN G P; ANDRIANOV V M; PIRUZYAN E S AUTHOR ADDRESS: INST. MOL. GENET., ACAD. SCI. USSR, MOSCOW, USSR. JOURNAL: MOL GENET MIKROBIOL VIRUSOL 0 (7). 1989. 11-13. 1989

FULL JOURNAL NAME: Molekulyarnaya Genetika Mikrobiologiya i Virusologiya

CODEN: MGMVD

RECORD TYPE: Abstract LANGUAGE: RUSSIAN

ABSTRACT: The gene transfer into plants using the genetic engineering methods gives us the possibility to obtain transgeneric plants having acquired the new traits. Some bacterial genes can be used for this purpose. Obtaining of a transgeneric plant harbouring the cytokinin synthesis gene ipt (gene 4) from the T-DNA of Agrobacterium tumefaciens Ti-plasmid seems to be useful. The expression of tumor agrobacterial ipt gene in transformed plant cells interferes with the normal growth and regulation of the whole plant. The successful transfer of the cloned ipt gene from the recombinant plasmid pGV0319 into the tobacco plant using Agrobacterium vectors and succeeding regeneration of phenotypically normal transgenic plants are reported in the present paper.

12/3,AB/64 (Item 49 from file: 5) DIALOG(R)File 5:Biosis Previews(R) (c) 2002 BIOSIS. All rts. reserv.

06644386 BIOSIS NO.: 000087086563

ENDOGENOUS CYTOKININ-INDUCED PR-PROTEIN SYNTHESIS IN

NICOTIANA-TABACUM PLANTS

AUTHOR: ZAKHAR'EV V M; TASHPULATOV A SH; NURKIYANOVA K M; TAL'YANSKII M E; KAPLAN I B; ATABEKOV I G; SKRYABIN K G

AUTHOR ADDRESS: INST. MOL. BIOL., ACAD. SCI. USSR, MOSCOW, USSR.

JOURNAL: DOKL AKAD NAUK SSSR 301 (3). 1988. 743-745. 1988

FULL JOURNAL NAME: Doklady Akademii Nauk Sssr

CODEN: DANKA

RECORD TYPE: Abstract LANGUAGE: RUSSIAN

ABSTRACT: Transgenic N. tabacum plants expressing isopentenyl transferase were created. A diagram describing the construction of the pAShT24 plasmid was presented. It was shown that intensive cytokinin synthesis in transgenic plants stimulates the production of PR-proteins and, possibly, increases the resistance of these plants to tobacco mosaic virus. The results of the study make it possible to establish a new approach for obtaining transgenic plants resistant to virus infections. 1988

12/3,AB/65 (Item 50 from file: 5) DIALOG(R)File 5:Biosis Previews(R) (c) 2002 BIOSIS. All rts. reserv.

06163519 BIOSIS NO.: 000085126671

HORMONAL REGULATION OF ZEATIN-RIBOSIDE ACCUMULATION BY CULTURED TOBACCO CELLS

AUTHOR: HANSEN C E; MEINS F JR; AEBI R

AUTHOR ADDRESS: FRIEDRICH MIESCHER-INST., P.O. BOX 2543, CH-4002 BASEL, SWITZERLAND.

JOURNAL: PLANTA (BERL) 172 (4). 1987. 520-525. 1987

FULL JOURNAL NAME: PLANTA (Berlin)

CODEN: PLANA

RECORD TYPE: Abstract LANGUAGE: ENGLISH

ABSTRACT: Auxin (11 .mu.M .alpha.-naphthaleneacetic acid) and cytokinin (1.4 .mu.M kinetin) regulate cytokinin accumulation by cytokinin-requiring (C-) and cytokinin-autotrophic (C+) lines of Havana 425 tobacco (Nicotiana tabacum L.) tissues. No trans-zeatin riboside (ZR) (< 0.5 pmol .cntdot. g-1 fresh weight) was detected in six C- and nine C+ lines grown for 14 d on auxin + cytokinin and auxin medium, respectively. C+ lines, but not Clines accumulated ZR (1.9-51. p mol .cntdot. g-1 fresh weight) when incubated on hormone-free medium but both lines accumulated ZR when incubated on kinetin medium. Therefore, it appears that kinetin treatment can induce ZR accumulation and that this accumulation is blocked by auxin treatment. Similar effects were obtained with some lines of cells autotrophic for both auxin and cytokinin. Tobacco plants carrying the dominant Habituated leaf-1 allele (Hl-1) differ from wild-type plants in that leaf-derived tissues in culture exhibit a C+ phenotype. No differences in ZR content were found in C+ leaf tissues from H1-1/H1-1 plants and C+ tissues that arise epigenetically in

wild-type plants. This indicates that the H-1 allele does not act to induce overproduction of ZR. The Hl-1 allele is known to have oncogenic functions similar to the isopentenyl transferase (ipt) locus of the Ti plasmid. Although Hl-1/Hl-1 cells transformed with Ti plasmids defective at the ipt locus are tumorigenic and hormone-autotrophic in culture, they contain low levels of ZR typical of non-transformed Hl-1 Hl-1 cells. Therefore, the high levels of ZR characteristic of cells transformed with wild-type Ti plasmids are not necessary for expression of the tumor phenotype.

1987

12/3,AB/66 (Item 51 from file: 5) DIALOG(R)File 5:Biosis Previews(R) (c) 2002 BIOSIS. All rts. reserv.

06074284 BIOSIS NO.: 000085037433 TUMOR FORMATION AND MORPHOGENESIS ON DIFFERENT NICOTIANA-SP AND HYBRIDS INDUCED BY AGROBACTERIUM-TUMEFACIENS T DNA MUTANTS AUTHOR: NACMIAS B; UGOLINI S; RICCI M D; PELLEGRINI M G; BOGANI P; BETTINI P; INZE D; BUIATTI M AUTHOR ADDRESS: DEP. ANIMAL BIOL. GENETICS, UNIV. FLORENCE, ITALY. JOURNAL: DEV GENET 8 (2). 1987. 61-72. 1987 FULL JOURNAL NAME: Developmental Genetics CODEN: DGNTD

RECORD TYPE: Abstract LANGUAGE: ENGLISH

ABSTRACT: A series of experiments are presented that have been performed to observe the interactions between Agrobacterium tumefaciens strains mutated in the T-DNA genes involved in indoleacetic acid and cytokinin biosynthesis and several Nicotiana species and hybrids. Infections were induced on leaf cuttings of Nicotiana debneyi, N. knightiana, N. clevelandii, N. bigelovii var. bigelovii, N. bigelovii var quadrivalvis, N. glauca, N. langsdorffi, the amphidiploid tumorous hybrid N. glauca .times. N. langsdorffi, and a nontumorous mutant of it. The effect of deletions of the Ti plasmid varied according to plant genotype. Insertion mutants in iaaM and iaaH suppressed tumor formation in N. langsdorffii, reduced it in N. bigelovii var quadrivalvis, had no effect in N. glauca and the two amphidiploid hybrids, and promoted tumorigenesis when compared to the wild-type Agrobacterium strain B6S3 in N. bigelovii var bigelovii, N. debneyi, and N. knightiana. The same mutations induced shoot formation in N. glauca, increased it in N. debneyi, and suppressed root formation in N. knightiana. On the other hand, an insertion mutation of the isopentenyl transferase gene (ipt-) had no effect in N. bigelovii var quadrivalvis, N. debneyi, the tumorous hybrid, suppressed tumor formation in N. langsdorffii, and inhibited it in N. glauca, the nontumorous hybrid, N. bigelovii var bigelovii, and N. knightiana. Insertion in ipt suppressed shoot formation in the nontumourous hybrid and inhibited it in the nontumorous amphidiploid and N. debneyi, while promoting root formation in N. glauca and N. debneyi. The suggestion of the existence of specific hormone equilibria necessary for the shift to each morphogenetic pattern was supported by experiments with exogenous hormone treatments of three genotypes (N. glauca, N. langsdorffii, and the nontumourous N. glauca .times. N. langsdorffii).

1987

12/3,AB/67 (Item 52 from file: 5) DIALOG(R)File 5:Biosis Previews(R) (c) 2002 BIOSIS. All rts. reserv.

06063253 BIOSIS NO.: 000085026402

TWO AGROBACTERIUM-TUMEFACIENS GENES FOR CYTOKININ BIOSYNTHESIS TI PLASMID-CODED ISOPENTENYLTRANSFERASES ADAPTED FOR FUNCTION IN PROKARYOTIC OR EUKARYOTIC CELLS

AUTHOR: HEINEMEYER W; BUCHMANN I; TONGE D W; WINDASS J D; ALT-MOERBE J; WEILER E W; BOTZ T; SCHROEDER J

AUTHOR ADDRESS: INST. BIOL. II, UNIV. FREIBURG, SCHAENZLESTR. 1, D-7800 FREIBURG, FRG.

JOURNAL: MOL GEN GENET 210 (1). 1987. 156-164. 1987 FULL JOURNAL NAME: Molecular & General Genetics

CODEN: MGGEA

RECORD TYPE: Abstract LANGUAGE: ENGLISH

ABSTRACT: Tzs and ipt are two Ti plsmid genes coding for proteins with isopentenyltransferase (IPT) activity in vitro. We cloned both genes for protein expression in Escherichia coli and in Agrobacterium tumefaciens, and we investigated differences between the two genes by analysing the properties of the proteins in vitro and in vivo. In vitro, extracts with tzs or ipt-coded proteins had high IPT activity, and the enzymes were identical in most properties. The most important difference was detected in vivo: the tzs-encoded protein was very active in cytokinin production, while the ipt protein required overexpression in order to obtain measurable activity in bacteria. In both cases, trans-zeatin was the major product of the gene activity. Formation of this cytokinin requires a hydroxylase function in addition to the IPT reaction. No such activity could be ascribed to tzs or ipt-encoded proteins in vitro or in vivo, but cytokinin hydroxylase activity was detected in cells and extracts of E. coli regardless of the presence or absence of the cytokinin genes. Based on these results it is proposed that both genes code for a single enzyme activity (isopentenyltransferase), that the genes and the proteins are adapted for function either in bacteria (tzs) or in transformed plant cells (ipt), and that in both prokaryotic and eukaryotic cells hydroxylation to trans-zeatin is a function contributed by host enzymes.

1987

12/3,AB/68 (Item 53 from file: 5) DIALOG(R)File 5:Biosis Previews (R) (c) 2002 BIOSIS. All rts. reserv.

BIOSIS NO.: 000084112513

INITIATION OF AUXIN AUTONOMY IN NICOTIANA-GLUTINOSA CELLS BY THE CYTOKININ-BIOSYNTHESIS GENE FROM AGROBACTERIUM-TUMEFACIENS AUTHOR: BINNS A N; LABRIOLA J; BLACK R C

AUTHOR ADDRESS: DEP. BIOL., UNIV. PENNSYLVANIA, PHILADELPHIA, PA. 19104-6018.

JOURNAL: PLANTA (BERL) 171 (4). 1987. 539-548. 1987

FULL JOURNAL NAME: PLANTA (Berlin)

CODEN: PLANA

RECORD TYPE: Abstract LANGUAGE: ENGLISH

ABSTRACT: Agrobacteria carrying mutations at the auxin-biosynthesizing loci (iaaH and iaaM of the Ti plasmid) induce shoot-forming tumors on many plant species. In some cases, e.g. Nicotiana glutinosa L., tumors induced by such mutant strains exhibit an unorganized and fully autonomous phenotype. These characteristics are stable in culture at both the tissue and cellular level. We demonstrate that the cytokinin -biosynthesis gene (ipt) of the Ti plasmid is responsible for the

induction of both auxin and cytokinin autonomy in N. glutinosa. Cloned cell lines carrying an ipt gene but lacking iaaH and iaaM are capable of accumulating indole-3-acetic acid. Interestingly, non-transformed N. glutinosa tissues exhibit an auxin-requiring phenotype when they are grown on medium supplemented with an exogenous supply of cytokinin. These results strongly indicate that exogenously supplied cytokinin does not mimic all the effects of the expression of the ipt gene in causing the auxin-autonomous growth of N. glutinosa cells.

1987

12/3,AB/69 (Item 54 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
(c) 2002 BIOSIS. All rts. reserv.

04745980 BIOSIS NO.: 000080049107
TUMOR GENES IN **PLANTS** T DNA ENCODED **CYTOKININ** BIOSYNTHESIS
AUTHOR: BUCHMANN I; MARNER F-J; SCHROEDER G; WAFFENSCHMIDT S; SCHROEDER J
AUTHOR ADDRESS: INST. FUER BIOL. II, UNIV. FREIBURG, SCHAENZLESTR. 1,
D-7800 FREIBURG, FRG.
JOURNAL: EMBO (EUR MOL BIOL ORGAN) J 4 (4). 1985. 853-860. **1985**FULL JOURNAL NAME: EMBO (European Molecular Biology Organization) Journal
CODEN: EMJOD

RECORD TYPE: Abstract LANGUAGE: ENGLISH

ABSTRACT: Whether gene 4 from the T-region of the Agrobacterium tumefaciens Ti plasmid codes for an enzyme of hormone biosynthesis was investigated. In a 1st set of experiments, gene 4 from octopine plasmid pTiAch5 and nopaline plasmid pTiC58 was expressed in Escherichia coli, and the gene products were identified by reaction with antiserum raised against a decapeptide derived from the DNA sequence of the gene. Extracts from cells expressing the gene contained high isopentenyl transferase activity catalyzing the formation of N6-(.DELTA.2isopentenyl) adenosine from 5'-AMP and .DELTA.2-isopentenylpyrophosphate. The cytokinin was identified by sequential high performance liquid chromatography and mass spectrometry. In a 2nd set of experiments it was shown that crown gall cells contained isopentenyl transferase activity and a protein of MW 27,000, which was identified as the product of gene 4 by reaction with the antiserum. Isopentenyl transferase activity was specifically inhibited by the antiserum. No comparable enzyme activity or immunoreactive protein was detected in cytokinin-autotrophic, T-DNA free tobacco cells. The results establish that gene 4 from the T-region of octopine and nopaline Ti plasmids codes for an enzyme of cytokinin biosynthesis.

1985

12/3,AB/70 (Item 1 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
(c) 2002 Inst for Sci Info. All rts. reserv.

07302954 Genuine Article#: 147XB Number of References: 11 Title: Rice transformation with a senescence-inhibition chimeric gene (ABSTRACT AVAILABLE)

Author(s): Fu YC; Ding YY; Liu XF; Sun CQ; Cao SY; Wang DJ; He SJ; Wang XK; Li LC; Tian WZ (REPRINT)

Corporate Source: CHINESE ACAD SCI, GENET INST/BEIJING 100101//PEOPLES R CHINA/ (REPRINT); CHINESE ACAD SCI, GENET INST/BEIJING 100101//PEOPLES R CHINA/; CHINA AGR UNIV,/BEIJING 100094//PEOPLES R CHINA/

Journal: CHINESE SCIENCE BULLETIN, 1998, V43, N21 (NOV), P1810-1815

ISSN: 1001-6538 Publication date: 19981100

Publisher: SCIENCE CHINA PRESS, 16 DONGHUANGCHENGGEN NORTH ST, BEIJING 100717, PEOPLES R CHINA

Language: English Document Type: ARTICLE

Abstract: A senescence-inhibition chimeric gene containing the specific promoter of SAG(12) and IPT gene was transferred into rice with the biolistic method. Results of PCR, Dot blotting and Southern blotting indicated that the chimeric gene had been integrated into rice genome. Analyses of GUS activity and cytokinin content in transgenic plants of rice and the observation of T-1 generation plant at grain formation stage indicated that the foreign gene was expressed.

12/3,AB/71 (Item 2 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
(c) 2002 Inst for Sci Info. All rts. reserv.

07048099 Genuine Article#: 118EM Number of References: 48
Title: Expression of the yeast mevalonate kinase gene in trangenic tobacco (ABSTRACT AVAILABLE)

Author(s): Champenoy S; Tourte M (REPRINT)

Corporate Source: IBMIG, UPRES 1221, LAB BIOL CELLULAIRE VEGETALE, 40 AVE RECTEUR PINEAU/F-86022 POITIERS//FRANCE/ (REPRINT); IBMIG, UPRES 1221, LAB BIOL CELLULAIRE VEGETALE/F-86022 POITIERS//FRANCE/

Journal: MOLECULAR BREEDING, 1998, V4, N4, P291-300

ISSN: 1380-3743 Publication date: 19980000

Publisher: KLUWER ACADEMIC PUBL, SPUIBOULEVARD 50, PO BOX 17, 3300 AA DORDRECHT, NETHERLANDS

Language: English Document Type: ARTICLE

Abstract: The coding region of the yeast mevalonate kinase gene (ERG12), under the control of the cauliflower mosaic virus (CaMV) 35S promoter, has been inserted in tobacco (Nicotiana tabacum cv. Paraguay Bell) using an Agrobacterium tumefaciens binary vector system. Integration and expression of the ERG12 chimaeric gene was demonstrated in several independent transformants in which specific mevalonate kinase (MK) activity in young plantlets was increased by about 60% on average. The expression of this MK gene was accompanied by phenotypical modifications, such as acceleration of regenerating processes, lateral bud growth, and peculiar flowering behaviour. A higher chlorophyll content all along the plant development, paralleled by an unusual starch accumulation in the leaves of young plantlets and, later, in roots of full-grown plants, was also detected. Overexpression of the MK gene led also to a stronger inhibition of cytokinin-induced plant growth by methyl jasmonate in transgenic plants. All these events may be interpreted as a possible modification of the hormonal balance in transgenic tobaccos.

12/3,AB/72 (Item 3 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
(c) 2002 Inst for Sci Info. All rts. reserv.

06866764 Genuine Article#: ZX908 Number of References: 59
Title: Two genes with similarity to bacterial response regulators are rapidly and specifically induced by cytokinin in Arabidopsis (ABSTRACT AVAILABLE)

Author(s): Brandstatter I; Kieber JJ (REPRINT)

Corporate Source: UNIV ILLINOIS, DEPT BIOL SCI, MOL BIOL

LAB/CHICAGO//IL/60607 (REPRINT); UNIV ILLINOIS, DEPT BIOL SCI, MOL BIOL LAB/CHICAGO//IL/60607

Journal: PLANT CELL, **1998**, V10, N6 (JUN), P1009-1019

ISSN: 1040-4651 Publication date: 19980600

Publisher: AMER SOC PLANT PHYSIOLOGISTS, 15501 MONONA DRIVE, ROCKVILLE, MD

20855

Language: English Document Type: ARTICLE

Abstract: Cytokinins are central regulators of plant growth and development, but little is known about their mode of action. By using differential display, we identified a gene, IBC6 (for induced by cytokinin), from etiolated Arabidopsis seedlings, that is induced rapidly by cytokinin. The steady state level of IBC6 mRNA was elevated within 10 min by the exogenous application of cytokinin, and this induction did not require de novo protein synthesis. IBC6 was not induced by other plant hormones or by light. A second Arabidopsis gene with a sequence highly similar to IBC6 was identified. This IBC7 gene also was induced by cytokinin, although with somewhat slower kinetics and to a lesser extent. The pattern of expression of the two genes was similar, with higher expression in leaves, rachises, and flowers and lower transcript levels in roots and siliques. Sequence analysis revealed that IBC6 and IBC7 are similar to the receiver domain of bacterial two-component response regulators. This homology, coupled with previously published work on the CKI1 histidine kinase homolog, suggests that these proteins may play a role in early cytokinin signaling.

12/3,AB/73 (Item 4 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
(c) 2002 Inst for Sci Info. All rts. reserv.

06398998 Genuine Article#: YP814 Number of References: 248
Title: The molecular basis of cytokinin action (ABSTRACT AVAILABLE)

Author(s): Hare PD (REPRINT); vanStaden J

Corporate Source: UNIV NATAL, DEPT BOT, RES UNIT PLANT GROWTH & DEV, PRIVATE BAG X01/ZA-3209 SCOTTSVILLE//SOUTH AFRICA/ (REPRINT)

Journal: PLANT GROWTH REGULATION, 1997, V23, N1-2 (OCT), P41-78

ISSN: 0167-6903 Publication date: 19971000

Publisher: KLUWER ACADEMIC PUBL, SPUIBOULEVARD 50, PO BOX 17, 3300 AA DORDRECHT, NETHERLANDS

Language: English Document Type: REVIEW

Abstract: Current understanding of cytokinin (CK) physiology at the cellular level results largely from the manipulation of endogenous CK levels by either application of exogenous CKs or the expression of CK biosynthetic transgenes, as well as the characterisation of single gene mutants. Cytokinins modulate changes in plant gene expression, which are in turn assumed to effect physiological and morphological changes with which CK action is associated. Presently, a major focus of investigation is elucidation of the biochemical events leading from the perception of CK to the manifestation of a response. Analysis of the expression patterns of CK-regulated genes and identification of their products provides one means of investigating CK action at the molecular level. Biochemical approaches have led to the identification of several soluble CK-binding proteins, although their functional roles in CK signalling largely remain uncertain. Conclusive identification of a bona fide CK receptor has yet to be achieved, although several potential candidates have been suggested. Pharmacological and molecular genetic strategies have implicated the involvement of signalling mechanisms likely to be involved in CK action. The apparent involvement of fluctuations in the concentration of intracellular Ca2+, changes in protein phosphorylation as well as DNA and/or protein methylation provide information concerning the types of proteins likely to be involved in the process. Dissection of CK signal transduction chains and elucidation of their interaction with other pathways that regulate plant growth and development is likely to be essential in understanding the mode of action of this poorly understood class of plant growth regulator.

However, integration of this knowledge with an improved understanding of the mechanisms whereby overall hormone homeostasis is regulated at the metabolic level will be necessary for comprehensive appreciation of the influence of CKs on **plant** morphology and physiology.

12/3,AB/74 (Item 5 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
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06284505 Genuine Article#: YF800 Number of References: 30
Title: The rms1 mutant of pea has elevated indole-3-acetic acid levels and reduced root-sap zeatin riboside content but increased branching controlled by graft-transmissible signal(s) (ABSTRACT AVAILABLE)
Author(s): Beveridge CA (REPRINT); Symons GM; Murfet IC; Ross JJ; Rameau C Corporate Source: INRA,GENET & AMELIORAT PLANTES STN, ROUTE ST CYR/F-78026
 VERSAILLES//FRANCE/ (REPRINT); UNIV TASMANIA,DEPT PLANT SCI/HOBART/TAS 7001/AUSTRALIA/; UNIV QUEENSLAND,DEPT BOT/BRISBANE/QLD 4072/AUSTRALIA/
Journal: PLANT PHYSIOLOGY, 1997, V115, N3 (NOV), P1251-1258
ISSN: 0032-0889 Publication date: 19971100
Publisher: AMER SOC PLANT PHYSIOLOGISTS, 15501 MONONA DRIVE, ROCKVILLE, MD 20855

Language: English Document Type: ARTICLE

Abstract: Rms1 is one of the series of five ramosus loci in pea (Pisum sativum L.) in which recessive mutant alleles confer increased branching at basal and aerial vegetative nodes. Shoots of the nonallelic rms1 and rms2 mutants are phenotypically similar in most respects. However, we found an up to 40-fold difference in root-sap zeatin riboside ([9R]Z) concentration between rms1 and rms2 plants. Compared with wild-type (WT) plants, the concentration of [9R]Z in rms1 root sap was very low and the concentration in rms2 root sap was slightly elevated. To our knowledge, the rms1 mutant is therefore the second ramosus mutant (rms4 being the first) to be characterized with low root-sap [9R]Z content. Like rms2, the apical bud and upper nodes of rms1 plants contain elevated indole-3-acetic acid levels compared with WT shoots. Therefore, the rms1 mutant demonstrates that high shoot auxin levels and low root-sap cytokinin levels are not necessarily correlated with increased apical dominance in pea. A graft-transmissible basis of action has been demonstrated for both mutants from reciprocal grafts between mutant and WT plants. Branching was also largely inhibited in rms1 shoots when grafted to rms2 rootstocks, but was not inhibited in rms2 shoots grafted to rms1 rootstocks. These grafting results are discussed, along with the conclusion that hormone-like signals other than auxin and cytokinin are also involved.

12/3,AB/75 (Item 6 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
(c) 2002 Inst for Sci Info. All rts. reserv.

06177862 Genuine Article#: YA111 Number of References: 35
Title: Increased content of endogenous cytokinins does riot delay
degradation of photosynthetic apparatus in tobacco (ABSTRACT AVAILABLE

Author(s): Synkova H (REPRINT); VanLoven K; Valcke R
Corporate Source: ACAD SCI CZECH REPUBL, INST EXPT BOT, KARLOVCE 1A/CZ-16000
PRAGUE 6//CZECH REPUBLIC/ (REPRINT); LIMBURGS UNIV CTR, DEPT SBG/B-3590
DIEPENBEEK//BELGIUM/

Journal: PHOTOSYNTHETICA, 1997, V33, N3-4, P595-608

ISSN: 0300-3604 Publication date: 19970000

Publisher: INST EXPERIMENTAL BOTANY, ACAD SCI CZECH REPUBLIC, NA KARLOVCE 1A, PRAGUE 6, CZECH REPUBLIC CS-160 00

Language: English Document Type: ARTICLE

Abstract: The effect of stress (long-term darkening) on the structure and functioning of the photosynthetic apparatus was studied in leaves of non-transformed as well as two types of ipt-transformed tobacco (Nicotiana tabacum cv. Petit Havana SR1) plants, The ipt -gene controlling the biosynthesis of cytokinins (CKs) was coupled to the light-inducible Pssu-promoter of Pisum sativum of to the heat-inducible hsp-promoter of Drosophila melanogaster. Pssu-ipt transgenic grafts with high contents of endogenous CKs retained their chlorophyll (Chi) content during a 15 d dark treatment while the SR1and heat-treated Phsp 70-ipt seedlings, which did not differ significantly in CKs content, lost up to 60 % of their Chi. The normalised variable fluorescence ratio (F-v/F-m) and oxygen evolution decreased dramatically in the course of continuous dark treatment, indicating a degradation of photosystem. 2 irrespective of the plant type. Changes in the polypeptide composition of thylakoid membranes, as analysed by SDS-PAGE, confirmed this degradation process. Light and electron microscopic observations of leaf sections, and of the ultrastructure of plastids showed changes corresponding to a degradation of the photosynthetic apparatus.

12/3,AB/76 (Item 7 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
(c) 2002 Inst for Sci Info. All rts. reserv.

06031915 Genuine Article#: XQ731 Number of References: 31
Title: Changes of both polypeptide pattern and sensitivity to
 cytokinin following transformation of periwinkle tissues with the
 isopentenyl transferase gene (ABSTRACT AVAILABLE)
Author(s): Carpin S; Garnier F; Andreu F; Chenieux JC; Rideau M (REPRINT);
 Hamdi S

Corporate Source: UNIV TOURS, LAB BIOL VEGETALE & BIOCHIM CELLULAIRE, EA 2106, 31 AVE MONGE/F-37200 TOURS//FRANCE/ (REPRINT); UNIV TOURS, LAB BIOL VEGETALE & BIOCHIM CELLULAIRE, EA 2106/F-37200 TOURS//FRANCE/

Journal: PLANT PHYSIOLOGY AND BIOCHEMISTRY, 1997, V35, N8 (AUG), P 603-609

ISSN: 0981-9428 Publication date: 19970800

Publisher: GAUTHIER-VILLARS, 120 BLVD SAINT-GERMAIN, 75280 PARIS CEDEX 06, FRANCE

Language: English Document Type: ARTICLE

Abstract: Two-dimensional gel electrophoresis was used to examine differences between the polypeptide patterns of an untransformed periwinkle callus line and a transformed line carrying the cytokinin biosynthesis gene isopentenyl transferase (ipt) under control of a light-inducible promoter. Both lines were cultured for three weeks on an auxin free medium with or without exogenously-added zeatin, in continuous light or in complete darkness. Firstly, it was found that exogenous cytokinin treatment increased the amount of at least 24 polypeptides and decreased the amount of three polypeptides in the untransformed line. Secondly, a marked decrease in the number and the amount of the polypeptides was observed in the 2D-gels from the transgenic line. Traces of two cytokinin up-regulated polypeptides, the amounts of which have been previously found to be correlated with the accumulation of indole alkaloids in periwinkle cells in vitro were present in this line. Lastly, exogenous cytokinin treatment had very little effect on the polypeptide pattern of the transgenic line. These data show that endogenously-produced cytokinin does not mimic the effect of exogenously-applied cytokinin on the polypeptide accumulation in periwinkle callus cultures, and that the ipt-transgenic line has become insensitive to exogenous cytokinin treatment.

DIALOG(R)File 34:SciSearch(R) Cited Ref Sci (c) 2002 Inst for Sci Info. All rts. reserv.

05899416 Genuine Article#: XF276 Number of References: 57
Title: Cytokinin/auxin control of apical dominance in Ipomoea nil
ABSTRACT AVAILABLE)

Author(s): Cline M (REPRINT) ; Wessel T; Iwamura H

Corporate Source: OHIO STATE UNIV, DEPT PLANT BIOL, 1735 NEIL AVE/COLUMBUS//OH/43210 (REPRINT); KYOTO UNIV, FAC AGR, DEPT AGR CHEM/KYOTO 606//JAPAN/

Journal: PLANT AND CELL PHYSIOLOGY, 1997, V38, N6 (JUN), P659-667

ISSN: 0032-0781 Publication date: 19970600

Publisher: JAPANESE SOC PLANT PHYSIOLOGISTS, SHIMOTACHIURI OGAWA HIGASHI KAMIKYOKU, KYOTO 602, JAPAN

Language: English Document Type: ARTICLE

Abstract: Although the concept of apical dominance control by the ratio of cytokinin to auxin is not new, recent experimentation with transgenic plants has given this concept renewed attention, In the present study, it has been demonstrated that cytokinin treatments can partially reverse the inhibitory effect of auxin on lateral bud outgrowth in intact shoots of Ipomoea nil, Although less conclusive, this also appeared to occur in buds of isolated nodes. Auxin inhibited lateral bud outgrowth when applied either to the top of the stump of the decapitated shoot or directly to the bud itself, However, the fact that cytokinin promotive effects on bud outgrowth are known to occur when cytokinin is applied directly to the bud suggests different transport tissues and/or sites of action for the two hormones, Cytokinin antagonists were shown in some experiments to have a synergistic effect with benzyladenine on the promotion of bud outgrowth. If the ratio of cytokinin to auxin does control apical dominance, then the next critical question is how do these hormones interact in this correlative process? The hypothesis that shoot-derived auxin inhibits lateral bud outgrowth indirectly by depleting cytokinin content in the shoots via inhibition of its production in the roots was not supported in the present study which demonstrated that the repressibility of lateral bud outgrowth by auxin treatments at various positions on the shoot was not correlated with proximity to the roots but rather with proximity to the buds, Results also suggested that auxin in subtending mature leaves as well as that in the shoot apex and adjacent small leaves may contribute to the apical dominance of a shoot.

12/3,AB/78 (Item 9 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
(c) 2002 Inst for Sci Info. All rts. reserv.

05867710 Genuine Article#: XD243 Number of References: 50
Title: Tobacco plants carrying a tms locus of Ti-plasmid origin and the Hl-1 allele are tumor prone (ABSTRACT AVAILABLE)
Author(s): Meyer AD; Aebi R; Meins F (REPRINT)
Corporate Source: FRIEDRICH MIESCHER INST, BOX 2543/CH-4002
BASEL//SWITZERLAND/ (REPRINT); FRIEDRICH MIESCHER INST,/CH-4002
BASEL//SWITZERLAND/

Journal: DIFFERENTIATION, 1997, V61, N4 (MAY), P213-221

ISSN: 0301-4681 Publication date: 19970500

Publisher: SPRINGER VERLAG, 175 FIFTH AVE, NEW YORK, NY 10010

Language: English Document Type: ARTICLE

Abstract: The autonomous growth of plant tumor cells is believed to result from their persistent loss of the requirement for growth hormones such as auxin and cytokinin. The partially dominant gene Habituated leaf-1 (HI-1) regulates the requirement of cultures tissues of Havana 425 tobacco (Nicotiana tabacum L.) for cytokinins. The HI-1 allele can partially restore the tumor phenotype in tobacco cells

transformed with a Agrobacterium tumefaciens Ti plasmid defective in the isopentenyl transferase locus, which encodes a key enzyme in cytokinin biosynthesis and is required for neoplastic growth. To investigate the oncogenic function of HI-1, we transformed wild-type (hl-1/hl-1) and Hl-1/Hl-1 tobacco plants with the tms locus derived from the limited-host-range Ti plasmid pTiAg162. This locus encodes enzymes for biosynthesis of the auxin indole-3-acetic acid. Grafting tests and measurements of the hormone requirement of cultured explants show that wound-induced overgrowths arising in tms transformed Hl-1 plants are tumorous. While some wound-induced overgrowths also formed in hl-1/hl-1 transformants, these showed slight hormone-autotrophic growth and weak tumorigenicity in grafting tests, In addition, Hl-1/Hl-1 tms/tms plants, but not hl-1/hl-1 tms/tms plants, spontaneously developed rooty teratomatous overgrowths, showed flowering abnormalities, and formed calli at the base of the stem in young seedlings. Thus, Hl-1 tms plants exhibit a tumor-prone phenotype, and in this regard closely resemble tumor-prone hybrids that arise in certain interspecific crosses of Nicotiana species. Our results show that the interaction of just two genetic elements - the mutant Hl-1 allele of the tobacco host with tms genes of Ti plasmid origin - are sufficient for a tumor-prone phenotype.

12/3,AB/79 (Item 10 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
(c) 2002 Inst for Sci Info. All rts. reserv.

05754079 Genuine Article#: WV255 Number of References: 36
Title: Growth pattern, tuber formation and hormonal balance in in vitro potato plants carrying ipt gene (ABSTRACT AVAILABLE)
Author(s): Ivana M (REPRINT); Lidiya S; Milos O; Oksana Z; Tatyana K;
Josef E; Jaroslava O; Svetlana G; Yurii R; Nina A
Corporate Source: ACAD SCI CZECH REPUBL, DE MONTFORT UNIV, NORMAN BORLAUG

Corporate Source: ACAD SCI CZECH REPUBL, DE MONTFORT UNIV, NORMAN BORLAUG INST PLANT SCI, INST EXPT BOT, KE DVORU 15/PRAGUE 16600 6//CZECH REPUBLIC/ (REPRINT); RUSSIAN ACAD SCI, INST PLANT PHYSIOL/MOSCOW 127236//RUSSIA/; ACAD SCI CZECH REPUBL, INST PLANT MOL BIOL/CESKE BUDEJOVICE 38000//CZECH REPUBLIC/; INST CROP PROD, /PRAGUE 16106 6//CZECH REPUBLIC/

Journal: PLANT GROWTH REGULATION, 1997, V21, N1 (JAN), P27-36

ISSN: 0167-6903 Publication date: 19970100

Publisher: KLUWER ACADEMIC PUBL, SPUIBOULEVARD 50, PO BOX 17, 3300 AA DORDRECHT, NETHERLANDS

Language: English Document Type: ARTICLE

Abstract: Nodal cuttings of in vitro grown potato plants (Solanum tuberosum, cv. Miranda) were transformed by a vector plasmid carrying ipt gene of Agrobacterium tumefaciens, From the initial teratoma stage 5 clones of transgenic plants (1, 2, 11, 13 and 15) were obtained, which displayed in varying degree shortening of the internodes, decrease of the leaf size, decrease of apical dominance and poor rooting. In addition, two of the clones (11 and 13) showed increased stolen and tuber formation. In all these clones the endogenous level of free cytokinins (CKs) was increased: from 40% in clone 11 to almost 300% in clone 1. Also free indole-3-acetic acid (IAA) level was increased, but to a lower degree; the maximal increase was 160% (clone 13), Applied kinetin or IAA (1 mg.l(-1)) strongly suppressed root and tuber formation in clones 11 and 13, although they did not affect or even stimulated these processes in control plants. For control plants the minimal medium sucrose concentration necessary for tuber initiation was 6% whereas in clone 1 1 plants 2% was sufficient. Different distribution of endogenous CKs and IAA was observed in clone 11 and control plants. The highest CK content was found in transgenic plants in stems and in controls in leaves. In clone 11 plants abscisic acid (ABA) level was significantly increased in comparison to the control throughout the

cultivation period. Ethylene formation was strongly increased the first week after the subcultivation and later on the difference between transgenic and control plants rapidly diminished. Reactions of clone 11 plants to red (RL) and blue light (BL) were similar to reactions of control plants. In RL clone 11 plants were tall and thin with stunted leaves; in BL they had a teratoma-like appearance and formed a very high number of tubers. The role of hormones in these changes in growth and tuber formation is discussed.

12/3,AB/80 (Item 11 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
(c) 2002 Inst for Sci Info. All rts. reserv.

05669711 Genuine Article#: WP334 Number of References: 36 Title: Selection of marker-free transgenic **plants** using the **isopentenyl transferase** gene (ABSTRACT AVAILABLE)

Author(s): Ebinuma H (REPRINT); Sugita K; Matsunaga E; Yamakado M Corporate Source: NIPPON PAPER IND CO LTD, CENT RES LAB, KITA KU, 5-21-1 OJI/TOKYO 114//JAPAN/ (REPRINT)

Journal: PROCEEDINGS OF THE NATIONAL ACADEMY OF SCIENCES OF THE UNITED STATES OF AMERICA, 1997, V94, N6 (MAR 18), P2117-2121

ISSN: 0027-8424 Publication date: 19970318

Publisher: NATL ACAD SCIENCES, 2101 CONSTITUTION AVE NW, WASHINGTON, DC 20418

Language: English Document Type: ARTICLE

Abstract: We have developed a new plant vector system for repeated transformation (called MAT for multi-auto-transformation) in which a chimeric ipt gene, inserted into the transposable element Ac, is used as a selectable marker for transformation, Selectable marker genes conferring antibiotic or herbicide resistance, used to introduce economically valuable genes into crop plants, have three major problems: (i) the selective agents have negative effects on proliferation and differentiation of plant cells; (ii) there is uncertainty regarding the environmental impact of many selectable marker genes; (iii) it is difficult to perform recurrent transformations using the same selectable marker to pyramid desirable genes, The MAT vector system containing the ipt gene and the Ac element is designed to overcome these difficulties, When tobacco leaf segments were transformed and selected, subsequent excision of the modified Ac produced marker-free transgenic tobacco plants without sexual crosses or seed production, In addition, the chimeric ipt gene could be visually used as a selectable marker for transformation of hybrid aspen (Populus sieboldii x Populus grandidentata). The chimeric ipt gene, therefore, is an attractive alternative to the most widely used selectable marker genes. The MAT vector system provides a promising way to shorten breeding time for genetically engineered crops, This method could be particularly valuable for fruit and forest trees, for which long generation times are a more significant barrier to breeding and genetic analysis.

12/3,AB/81 (Item 12 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
(c) 2002 Inst for Sci Info. All rts. reserv.

05591252 Genuine Article#: WJ343 Number of References: 45 Title: Sterols and polyamines in IPT-transformed tobacco plants (ABSTRACT AVAILABLE)

Author(s): Geuns JMC (REPRINT); VanLoenhout HEM; Valcke RLM; VanLoven K; Redig P; Veselov SY; Kudoyarova GR; VanOnckelen HA; Vendrig JC Corporate Source: KATHOLIEKE UNIV LEUVEN, LAB PLANT PHYSIOL, MERECIERLAAN 92/B-3001 HEVERLEE//BELGIUM/ (REPRINT); LIMBURGS UNIV CTR, DEPT SBG/B-3590 DIEPENBEEK//BELGIUM/; UNIV INSTELLING ANTWERP, LAB PLANT

PHYSIOL/B-2610 WILRIJK//BELGIUM/; RUSSIAN ACAD SCI, INST BIOL, BASHKIR SCI CTR/UFA 450054/BASHKORTOSTAN/RUSSIA/

Journal: PHYTOCHEMISTRY, 1997, V44, N5 (MAR), P797-804

ISSN: 0031-9422 Publication date: 19970300

Publisher: PERGAMON-ELSEVIER SCIENCE LTD, THE BOULEVARD, LANGFORD LANE,

KIDLINGTON, OXFORD, ENGLAND OX5 1GB

Language: English Document Type: ARTICLE

Abstract: Free sterol and free polyamine contents were determined in the apex and the leaves of control and Pssu-ipt transformed tobacco (Nicotiana tabacum L. cv. Petit Havana SR1). The older leaves of ipt-transformed plants contained a much higher putrescine (put) content than those of control SR1 plants, whereas no significant differences for spermidine (spd) or spermine (spm) were found between control and ipt plants. Putrescine content corresponded well with endogenous cytokinin (free-bases) content and with ornithine- and ornithine-decarboxylose (ODC and ADC) activities. Plants transformed with ipt were characterized by a higher sterol content in the leaves and by a delay in the increase in the stigmasterol/sitosterol ratio that occurs from the upper to the

12/3,AB/82 (Item 13 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
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05572934 Genuine Article#: WH449 Number of References: 31
Title: Cultivars of hexaploid wheat of contrasting stature and chlorophyll retention differ in cytokinin content and responsiveness (
ABSTRACT AVAILABLE)

Author(s): Banowetz GM (REPRINT)

Corporate Source: USDA ARS, 3450 SW CAMPUS WAY/CORVALLIS//OR/97331 (REPRINT)

Journal: ANNALS OF BOTANY, 1997, V79, N2 (FEB), P185-190

lower leaves. Copyright (C) 1997 Elsevier Science Ltd.

ISSN: 0305-7364 Publication date: 19970200

Publisher: ACADEMIC PRESS LTD, 24-28 OVAL RD, LONDON, ENGLAND NW1 7DX

Language: English Document Type: ARTICLE

Abstract: The work reported here compared cytokinin content and sensitivity in a selection of hexaploid wheat (Triticum aestivum L.) cultivars using the following measurements: leaf cytokinins at three time points during light-growth and at four 24 h intervals after light-grown plants were transferred to darkness; sensitivity of root growth to direct applications of isopentenyl adenosine ([9R]iP); and, sensitivity of germination and subsequent root and shoot growth to 18 h imbibition of seeds in benzyladenine (BA).

Accumulation of zeatin riboside-type cultivars was greatest during light-growth in Tibet Dwarf a wheat with an extreme dwarf phenotype, intermediate in Omar standard and dwarf cultivars, and lowest in the standard and dwarf versions of Itana. Cytokinin levels were otherwise not directly correlated to plant stature in these wheats. There were no cultivar-associated qualitative differences in the types of ${\it cytokinins}$ detected in this study. During the 16 h light period, the content of zeatin riboside-type cytokinins increased up to tenfold and then declined to basal levels during dark growth. Chlorophyll retention during dark-growth was correlated with leaf cytokinin content. Data collected at a restricted number of sampling points during dark-growth suggested a cyclic accumulation of [9R] iP-type cytokinins and the apparent cycle in Tibet Dwarf was offset by 24 h. Tibet Dwarf showed the greatest root growth inhibition after exposure of seedling roots to [9R]iP or imbibition of seeds in BA. Neither of these treatments affected shoot growth in any of the cultivars. (C) 1997 Annals of Botany Company.

12/3,AB/83 (Item 14 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
(c) 2002 Inst for Sci Info. All rts. reserv.

04995885 Genuine Article#: UY013 Number of References: 71
Title: CYTOKININS IN PLANT SENESCENCE - FROM SPRAY AND PRAY TO
CLONE AND PLAY (Abstract Available)

Author(s): GAN SS; AMASINO RM

Corporate Source: UNIV WISCONSIN, DEPT BIOCHEM, 420 HENRY

MALL/MADISON//WI/53706; UNIV WISCONSIN, DEPT BIOCHEM/MADISON//WI/53706

Journal: BIOESSAYS, 1996, V18, N7 (JUL), P557-565

ISSN: 0265-9247

Language: ENGLISH Document Type: REVIEW

Abstract: Three approaches have been used to investigate the inhibitory role of the cytokinin class of phytohormones in plant senescence: external application of cytokinins, measurement of endogenous cytokinin levels before and during senescence, and manipulation of endogenous cytokinin production in transgenic plants. In transgenic plant studies, endogenous cytokinin levels are manipulated by expression of IPT, a gene encoding isopentenyl transferase. Transgenic plants expressing IPT from a variety of promoters exhibit developmental and morphological alterations and often display retarded leaf senescence. A recently developed autoregulatory senescence-inhibition system targets cytokinin production quantitatively, spatially and temporally, and results in transgenic plants that exhibit significantly delayed senescence without abnormalities. These transgenic studies not only confirm the regulatory role of cytokinins in plant senescence, but also provide a way to manipulate senescence for potential agricultural applications.

12/3,AB/84 (Item 15 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
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04597152 Genuine Article#: TW208 Number of References: 46
Title: TRANSFER-RNA IS THE SOURCE OF EXTRACELLULAR ISOPENTENYLADENINE IN A
TI-PLASMIDLESS STRAIN OF AGROBACTERIUM-TUMEFACIENS (Abstract Available
)

Author(s): GRAY J; GELVIN SB; MEILAN R; MORRIS RO
Corporate Source: UNIV MISSOURI, DEPT AGRON/COLUMBIA//MO/65211; UNIV
MISSOURI, DEPT BIOCHEM/COLUMBIA//MO/65211; PURDUE UNIV, DEPT BIOL SCI/W
LAFAYETTE//IN/47909

Journal: PLANT PHYSIOLOGY, **1996**, V110, N2 (FEB), P431-438 ISSN: 0032-0889

Language: ENGLISH Document Type: ARTICLE

Abstract: Even in the absence of the classical Ti plasmid-encoded cytokinin biosynthetic genes ipt and tzs, Agrobacterium tumefaciens strains still release significant amounts of the cytokinin isopentenyladenine (iP) into the culture medium (R.W. Kaiss-Chapman and R.O. Morris [1977] Biochem Biophys Res Commun 76: 453-459). A potential source of the iP is isopentenylated transfer RNA (tRNA), which, in turn, is synthesized by the activity of tRNA:isopentenyltransferase encoded by the bacterial miaA gene. To determine whether secreted iP had its origin in isopentenylated tRNA, a miaA(-) deletion/insertion mutant was prepared and reconstructed in Agrobacterium tumefaciens in vivo. The mutant no longer possessed tRNA:isopentenylation activity and no longer released iP into the extracellular medium. Transfer RNA therefore makes a small but significant contribution to the total amount of cytokinin normally secreted by Agrobacterium strains. tRNA-mediated synthesis may also account for cytokinin production by other plant -associated bacteria, such as Rhizobia, that have been reported to

secrete similarly low levels of nonhydroxylated cytokinins.

12/3,AB/85 (Item 16 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
(c) 2002 Inst for Sci Info. All rts. reserv.

04549569 Genuine Article#: TR821 Number of References: 10
Title: TISSUE-CULTURE AND TRANSFORMATION OF OENOTHERA-BIENNIS (Abstract Available)

Author(s): PAVINGEROVA D; GALIS I; ONDREJ M

Corporate Source: ACAD SCI CZECH REPUBL, INST PLANT MOLEC BIOL, BRANISOVSKA 31/CR-37005 CESKE BUDEJOVICE//CZECH REPUBLIC/

Journal: BIOLOGIA PLANTARUM, 1996, V38, N1, P27-32

ISSN: 0006-3134

Language: ENGLISH Document Type: ARTICLE

Abstract: Five cultivars of Oenothera biennis have been tested for callogenesis and organogenesis on different media. The cultivar CV3 has been transformed by Agrobacterium tumefaciens strain which introduces into the plant genome kanamycin resistance gene and the T-DNA ipt gene which causes increased levels of cytokinins.

Transformed tissues showed elevated levels of cytokinins and grew as teratomas forming clumps of short, branched shoots with small modified leaves. Roots appeared rarely in later subcultivations of some teratomous clones.

12/3,AB/86 (Item 17 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
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04482464 Genuine Article#: TG238 Number of References: 27
Title: THE EFFECT OF AN ELEVATED CYTOKININ LEVEL USING THE IPT
GENE AND N-6-BENZYLADENINE ON SINGLE NODE AND INTACT POTATO PLANT
TUBERIZATION IN-VITRO (Abstract Available)

Author(s): GALIS I; MACAS J; VLASAK J; ONDREJ M; VANONCKELEN HA
Corporate Source: ACAD SCI CZECH REPUBL, INST PLANT MOLEC BIOL, BRANISOVSKA
31/CR-37005 CESKE BUDEJOVICE//CZECH REPUBLIC/; UNIV INSTELLING
ANTWERP, DEPT BIOL/B-2610 WILRIJK//BELGIUM/

Journal: JOURNAL OF PLANT GROWTH REGULATION, 1995, V14, N3 (SUM), P 143-150

ISSN: 0721-7595

Language: ENGLISH Document Type: ARTICLE

Abstract: Two models of potato (Solanum tuberosum L.) tuberization in vitro (intact plants and single nodes) were used to study the role of cytokinins in this process. We applied hormone in two different ways. The exogenous addition of 10 mg . L(-1) N-6-benzyladenine (BA) into the tuberization medium resulted in advanced tuber formation in intact plants, and microtubers appeared 10-20 days earlier than in the experiments in which no cytokinin was supplied. Transformation with the Agrobacterium tumefaciens ipt gene provided potato clones with endogenously elevated cytokinin levels (3-20 times higher zeatin riboside content in different clones). The onset of tuberization in intact ipt-transformed plants with low transgene expression was advanced in comparison with control material, and exogenously applied BA further promoted the tuberization process. On the contrary, tuberization was strongly inhibited in ipt-transformed nodes, and an external increase of the cytokinin level caused complete inhibition of explant growth. In untransformed (control) nodes cytokinin application resulted in primary and secondary tuber formation, which depended on the BA concentration in cultivation media.

12/3,AB/87 (Item 18 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
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04206806 Genuine Article#: RN158 Number of References: 43
Title: PHENOTYPE MODIFICATION AND ENHANCED TOLERANCE TO INSECT PESTS BY
REGULATED EXPRESSION OF A CYTOKININ BIOSYNTHESIS GENE

Author(s): SMIGOCKI AC

Corporate Source: USDA ARS, PLANT MOLEC BIOL LAB/BELTSVILLE//MD/20705

Journal: HORTSCIENCE, 1995, V30, N5 (AUG), P967-969

ISSN: 0018-5345

Language: ENGLISH Document Type: ARTICLE

12/3,AB/88 (Item 19 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
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04168686 Genuine Article#: RK531 Number of References: 20
Title: EFFECT OF APEX EXCISION AND REPLACEMENT BY 1-NAPHTHYLACETIC ACID ON
CYTOKININ CONCENTRATION AND APICAL DOMINANCE IN PEA-PLANTS
(Abstract Available)

Author(s): LI CJ; GUEVARA E; HERRERA J; BANGERTH F
Corporate Source: UNIV HOHENHEIM, INST OBST GEMUSE & WEINBAU 370/D-70593
STUTTGART//GERMANY/; UNIV HOHENHEIM, INST OBST GEMUSE & WEINBAU
370/D-70593 STUTTGART//GERMANY/; UNIV COSTA RICA, FAC AGRON/SAN
JOSE//COSTA RICA/

Journal: PHYSIOLOGIA PLANTARUM, **1995**, V94, N3 (JUL), P465-469 ISSN: 0031-9317

Language: ENGLISH Document Type: ARTICLE

Abstract: As known from literature lateral buds from pea (Pisum sativum) plants are released from apical dominance when repeatedly treated with exogenous cytokinins. Little is known, however, about the endogenous role of cytokinins in this process and whether they interact with basipolar transported IAA, generally regarded as the main signal controlling apical dominance. This paper presents evidence that such an interaction exists.

The excision of the apex of pea plants resulted in the release of inhibited lateral buds from apical dominance (AD). This could be entirely prevented by applying 1-naphthylacetic acid (NAA) to the cut end of the shoot. Removal of the apex also resulted in a rapid and rather large increase in the endogenous concentrations of zeatin riboside (ZR), isopentenyladenosine (iAdo) and an as yet unidentified polar zeatin derivative in the node and internode below the point of decapitation. This accumulation of ZR and iAdo, was strongly reduced by the application of NAA. The observed increase in cytokinin concentration preceded the elongation of the lateral buds, suggesting that endogenous cytokinins play a significant role in the release of lateral buds from AD. However, the effect of NAA on the concentration of cytokinins clearly demonstrated the dominant role of the polar basipetally transported auxin in AD. The results suggest a mutual interaction between the basipolar IAA transport system and cytokinins obviously produced in the roots and transported via the xylem into the stem of the pea plants.

12/3,AB/89 (Item 20 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
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03950663 Genuine Article#: QU979 Number of References: 27
Title: APICAL DOMINANCE IN RHIZOMES OF QUACKGRASS, ELYTRIGIA REPENS - THE
EFFECT OF AUXIN, CYTOKININS, AND ABSCISIC-ACID (Abstract

Available)

Author(s): TAYLOR JS; ROBERTSON JM; HARKER KN; BHALLA MK; DALY EJ; PEARCE DW

Corporate Source: AGR CANADA, RES STN, BAG SERV 5000/LACOMBE/AB T0C1S0/CANADA/; UNIV CALGARY, DEPT BIOL SCI/CALGARY/AB T2N 1N4/CANADA/Journal: CANADIAN JOURNAL OF BOTANY-REVUE CANADIENNE DE BOTANIQUE, 1995, V73, N2 (FEB), P307-314

ISSN: 0008-4026

Language: ENGLISH Document Type: ARTICLE

Abstract: Experiments were designed to determine the impact of abscisic acid, indole-3-acetic acid, and cytokinins on dormancy of quackgrass (Elytrigia repens (L.) Nevski) rhizome axillary buds using exogenous hormone treatments and analysis of endogenous hormones. Exogenous hormone treatments were applied in solution or in lanolin paste to 5-node segments of rhizome with an apical tip intact or removed. Abscisic acid inhibited bud growth except at concentrations of 0.5-1 mu g ..mL(-1) when it stimulated growth: this appeared to be based on an inhibition of apical dominance rather than a stimulation of bud growth per se. Both indole-3-acetic acid and cytokinins stimulated bud growth, indole-3-acetic acid at concentrations of 0.5-5 mu g . mL(-1) and cytokinins at higher concentrations (i.e,, 10-100 mu g mL(-1)). Indole-3-acetic acid also increased elongation of the buds, whereas abscisic acid and cytokinins did not. Levels of endogenous hormones were measured in bud samples: indole-3-acetic acid was quantified as its methyl ester by combined gas chromatography - mass spectrometry - selected ion monitoring; abscisic acid was quantified as its methyl ester by gas chromatography - electron capture; and cytokinins were quantified using a soybean callus bioassay. Hormone levels were generally higher in the most active buds of a 5-node section. Abscisic acid was also measured in buds 24 h after sheath leaf removal, a practice known to promote bud sprouting. Sheath leaf removal had no significant effect on abscisic acid levels.

12/3,AB/90 (Item 21 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
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03572990 Genuine Article#: PN558 Number of References: 33
Title: INCREASED LEVEL OF CYTOKININ RIBOSIDES IN JASMONIC
ACID-TREATED POTATO (SOLANUM-TUBEROSUM) STEM NODE CULTURES (Abstract Available)

Author(s): DERMASTIA M; RAVNIKAR M; VILHAR B; KOVAC M Corporate Source: NATL INST BIOL, KARLOVSKA 19/LJUBLJANA 61000//SLOVENIA/ Journal: PHYSIOLOGIA PLANTARUM, 1994, V92, N2 (OCT), P241-246 ISSN: 0031-9317

Language: ENGLISH Document Type: ARTICLE

Abstract: Cytokinin free bases, ribosides and 9-glucosides were measured in stem node cultures of potato (Solanum tuberosum L. cv. Ulster Sceptre) in the presence or absence of 1 mu M jasmonic acid (JA) to examine whether or not their changed levels were part of the JA-induced growth response. The enhanced growth response in JA-treated plantlets included: expanded root systems, extended leaf areas, increased number of nodes, and enlarged stem diameters. The protein analysis revealed a substantial decrease in a 62-kDa polypeptide. On a dry weight basis, the levels of ribulose-1,5-biphosphate carboxylase/oxygenase (RuBP carboxylase, EC 4.1.1.39) and chlorophylls a and b were constant. The total concentration of endogenous cytokinins remained virtually the same in control and treated plantlets; but in JA-treated plantlets the amount of cytokinin free bases and cytokinin 9-glucosides decreased. In addition, the level of cytokinin ribosides was elevated. The ratio between active and inactive cytokinins increased from 1.2

to 2.1, which correlates with the enhanced growth of potato plantlets grown on 1 mu M JA. Thus the observed growth and developmental changes may be a consequence of the measured altered cytokinin level. However, significant morphological alterations of the potato plantlets treated with JA may also be a result of the changed critical cytokinin concentration or critical ratios of cytokinins to auxins and JA, rather than their absolute concentrations.

12/3,AB/91 (Item 22 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
(c) 2002 Inst for Sci Info. All rts. reserv.

03419064 Genuine Article#: NE203 Number of References: 156
Title: BACTERIAL GENES MODIFYING HORMONAL BALANCES IN **PLANTS** (
Abstract Available)

Author(s): GAUDIN V; VRAIN T; JOUANIN L

Corporate Source: INRA, BIOL CELLULAIRE LAB, ROUTE ST CYR/F-78026 VERSAILLES//FRANCE/; AGR CANADA, RES STN/VANCOUVER V6T 1X2/BC/CANADA/ Journal: PLANT PHYSIOLOGY AND BIOCHEMISTRY, 1994, V32, N1 (JAN-FEB)

, P11-29 ISSN: 0981-9428

Language: ENGLISH Document Type: REVIEW

Abstract: A number of microorganisms that interact with plants use the same hormonal signals as plants. Four phytopathogenic bacteria causing either ''olive knot'' disease, ''leafy galls'', ''crown gall'' or ''hairy root'' disease in plants are among those most studied. The last two species induce proliferation resembling normal tissue and organ formation. Crown gall and hairy root are due to the expression of oncogenes carried on a bacterial DNA fragment, which is transferred and integrated into the plant genome during infection. These oncogenes modify the plant hormonal balances or the hormone signal perception of the cells. In this review, we describe the different types of oncogenes present in several microorganisms, their functions when they are known, and die morphological, physiological, and developmental modifications that are induced when these oncogenes are introduced into plants.

12/3,AB/92 (Item 23 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
(c) 2002 Inst for Sci Info. All rts. reserv.

03268060 Genuine Article#: NT120 Number of References: 44
Title: ONCOGENE ARRANGEMENT IN A SHOOTY STRAIN OF AGROBACTERIUM-TUMEFACIENS
(Abstract Available)

Author(s): DREVET C; BRASILEIRO ACM; JOUANIN L

Corporate Source: INRA, BIOL CELLULAIRE LAB, ROUTE ST CYR/F-78026 VERSAILLES//FRANCE/; INRA, BIOL CELLULAIRE LAB/F-78026 VERSAILLES//FRANCE/

Journal: PLANT MOLECULAR BIOLOGY, 1994, V25, N1 (APR), P83-90

ISSN: 0167-4412

Language: ENGLISH Document Type: ARTICLE

Abstract: The Agrobacterium tumefaciens nopaline strain 82. 139 induces non-teratogenic shooty tumours on several plant species. We have determined the position of the T-region oncogenes in a 11.4 kb Xba I fragment which shows a general organization similar to its pTiC58 counterpart. Sequence analysis of the 4.7 kb right part of this fragment allowed us to identify the pTi82.139 ipt, 6b and nos coding sequences. pTi82.139 lacks the 6a gene, which lies between the ipt and 6b genes in pTiC58. The intervening region between the 6b and the nos genes contains an additional ORF with homology to ORF 21 (transcript 3') from the TR-DNA of octopine strain pTi15955.

12/3,AB/93 (Item 24 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
(c) 2002 Inst for Sci Info. All rts. reserv.

O3150931 Genuine Article#: NJ081 Number of References: 45
Title: THE FAS OPERON OF RHODOCOCCUS FASCIANS ENCODES NEW GENES REQUIRED FOR EFFICIENT FASCIATION OF HOST PLANTS (Abstract Available)
Author(s): CRESPI M; VEREECKE D; TEMMERMAN W; VANMONTAGU M; DESOMER J
Corporate Source: STATE UNIV GHENT, GENET LAB, KL LEDEGANCKSTR 35/B-9000
GHENT//BELGIUM/; STATE UNIV GHENT, GENET LAB/B-9000 GHENT//BELGIUM/
Journal: JOURNAL OF BACTERIOLOGY, 1994, V176, N9 (MAY), P2492-2501
ISSN: 0021-9193

Language: ENGLISH Document Type: ARTICLE

Abstract: Three virulence loci (fas, aft, and hgp) of Rhodococcus fascians D188 have been identified on a 200-kb conjugative linear plasmid (pFiD188). The fns locus was delimited to a 6.5-kb DNA fragment by insertion mutagenesis, single homologous disruptive recombination, and in trans complementation of different avirulent insertion mutants. The locus is arranged as a large operon containing six open reading frames whose expression is specifically induced during the interaction with host plants. One predicted protein is homologous to P-450 cytochromes from actinomycetes. The putative ferredoxin component is of a novel type containing additional domains homologous to transketolases from chemoautotrophic, photosynthetic, and methylotrophic microorganisms. Genetic analysis revealed that fas encodes, in addition to the previously identified ipt, at least two new genes that are involved in fasciation development, one of which is only required on older tobacco plants.

12/3,AB/94 (Item 25 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
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03077941 Genuine Article#: NA966 Number of References: 25
Title: BRANCHING MUTANT RMS-2 IN PISUM-SATIVUM - GRAFTING STUDIES AND
ENDOGENOUS INDOLE-3-ACETIC-ACID LEVELS (Abstract Available)
Author(s): BEVERIDGE CA; ROSS JJ; MURFET IC

Corporate Source: UNIV TASMANIA, DEPT PLANT SCI, GPO BOX 252C/HOBART/TAS 7001/AUSTRALIA/

Journal: PLANT PHYSIOLOGY, **1994**, V104, N3 (MAR), P953-959

ISSN: 0032-0889

Language: ENGLISH Document Type: ARTICLE

Abstract: Isogenic lines of pea (Pisum sativum L.) were used to determine the physiological site of action of the Rms-2 gene, which maintains apical dominance, and its effect on endogenous free indole-3-acetic acid (IAA) levels. In mutant rms-2 scions, which normally produce lateral branches below node 3 and above node 7, apical dominance was almost fully restored by grafting to Rms-2 (wildtype) stocks. In the reciprocal grafts, rms-2 stacks did not promote branching in wild-type shoots. Together, these results suggest that the Rms-2 gene inhibits branching in the shoot of pea by controlling the synthesis of a translocatable (hormone-like) substance that is produced in the roots and/or cotyledons and in the shoot. At all stages, including the stage at which aerial lateral buds commence outgrowth, the level of IAA in rms-a shoots was elevated (up to 5-fold) in comparison with that in wild-type shoots. The internode length of rms-2 plants was 40% less than in wild-type plants, and the mutant plants allocated significantly more dry weight to the shoot than to the root in comparison with wild-type plants. Crafting to wild-type stocks did not normalize IAA levels or internode length in rms-2 scions, even though it inhibited branching,

suggesting that the involvement of Rms-2 in the control of IAA level and internode length may be confined to processes in the shoot.

12/3,AB/95 (Item 26 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
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02704244 Genuine Article#: LX867 Number of References: 55
Title: APPLICATION OF CYTOKININS TO FLOWERS TO INCREASE POD SET IN
LUPINUS-ANGUSTIFOLIUS L (Abstract Available)

Author(s): ATKINS CA; PIGEAIRE A

Corporate Source: UNIV WESTERN AUSTRALIA, DEPT BOT/NEDLANDS/WA 6009/AUSTRALIA/; UNIV WESTERN AUSTRALIA, COOPERAT RES CTR LEGUMESMEDITERRANEAN AGR/NEDLANDS/WA 6009/AUSTRALIA/

Journal: AUSTRALIAN JOURNAL OF AGRICULTURAL RESEARCH, 1993, V44, N8, P1799-1819

ISSN: 0004-9409

Language: ENGLISH Document Type: ARTICLE

Abstract: Exogenous application of a 2 mol m-3 buffered solution of N6-benzylamninopurine (BAP) to flowers on the main stem inflorescence of Lupinus angustifolius L. cv. Danja profoundly altered reproductive development by reducing post-anthesis abscission of flowers and small pods. The same effect of BAP was recorded for a mutant (abs-) of cv. Danja, in which organ abscission was completely absent, indicating that localized application of cytokinin enhanced reproductive development rather than reduced pedicel abscission per se in the parent line. Application to pedicel and sepals at the open flower stage completely eliminated flower abortion on the main inflorescence, compared with less than 50% pod initiation on untreated inflorescences, more than doubled final pod yield on the main inflorescence and increased the number of mature pods on the whole plant by 33%. A single dose of BAP, to an inflorescence which bore flowers ranging in their stage of development from post-anthesis to immature flower buds, significantly increased the number of pods initiated and at final harvest, measured on a per plant basis. A number of synthetic and naturally occurring cytokinins, including zeatin riboside and dihydrozeatin riboside, were also effective. BAP application induced a longer period of flowering and resulted in a considerably thickened raceme. This was most marked at the distal end which showed enhanced cambial development, and secondary vascularization compared with untreated controls. The positive effects of BAP application on pod initiation were not restricted to cv. Danja but were found also for cv. Warrah and cv. Gungurru, both of which have enhanced pod set compared with Danja. Enhanced pod initiation on the main inflorescence generally reduced the number of pods developing on branch inflorescences. Additional application of BAP to flowers on branches, even at the most opportune time and at the most effective site, did not enhance pod initiation and, in some cases, significantly reduced pod set on these branches. The data indicate that it would be very difficult to exploit the positive effect of exogenous cytokinin application on pod set in field crops of lupin. However, selection or genetic engineering of plants with higher levels of endogenous cytokinins in flowers or flower parts at anthesis may provide a means by which to assess the importance of this factor in determining yield stability.

12/3,AB/96 (Item 27 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
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02507635 Genuine Article#: LG170 Number of References: 54
Title: HORMONAL CHARACTERIZATION OF TRANSGENIC TOBACCO PLANTS
EXPRESSING THE ROLC GENE OF AGROBACTERIUM-RHIZOGENES TL-DNA

Abstract Available)

Author(s): NILSSON O; MORITZ T; IMBAULT N; SANDBERG G; OLSSON O
Corporate Source: SWEDISH UNIV AGR SCI, DEPT FOREST GENET & PLANT
PHYSIOL/S-90183 UMEA//SWEDEN/; SWEDISH UNIV AGR SCI, DEPT FOREST GENET &
PLANT PHYSIOL/S-90183 UMEA//SWEDEN/; UMEA UNIV, DEPT PLANT
PHYSIOL/S-90187 UMEA//SWEDEN/

Journal: PLANT PHYSIOLOGY, 1993, V102, N2 (JUN), P363-371

ISSN: 0032-0889

Language: ENGLISH Document Type: ARTICLE

Abstract: Transgenic tobacco (Nicotiana tabacum L. cv Wisconsin 38) plants expressing the Agrobacterium rhizogenes rolC gene under the control of the cauliflower mosaic virus 35S RNA promoter were constructed. These plants displayed several morphological alterations reminiscent of changes in indole-3-acetic acid (IAA), cytokinin, and gibberellin (GA) content. However, investigations showed that neither the IAA pool size nor its rate of turnover were altered significantly in the rolC plants. The biggest difference between rolC and wild-type plants was in the concentrations of the cytokinin, isopentenyladenosine (iPA) and the gibberellin GA19. Radioimmunoassay and liquid chromatography-mass spectrometry measurements revealed a drastic reduction in rolC plants of iPA as well as in several other cytokinins tested, suggesting a possible reduction in the synthesis rate of cytokinins. Furthermore, gas chromatography-mass spectrometry quantifications of GA19 showed a 5- to 6-fold increase in rolC plants compared with wild-type plants, indicating a reduced activity of the GA19 oxidase, a proposed regulatory step in the gibberellin biosynthesis. Thus, we conclude that RolC activity in transgenic plants leads to major alterations in the metabolism of cytokinins and gibberellins.

12/3,AB/97 (Item 28 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
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02452864 Genuine Article#: LC487 Number of References: 36 Title: LEVELS AND LOCATION OF EXPRESSION OF THE

AGROBACTERIUM-TUMEFACIENS PTIA6 IPT GENE PROMOTER IN TRANSGENIC TOBACCO (Abstract Available)

Author(s): STRABALA TJ; CROWELL DN; AMASINO RM

Corporate Source: UNIV MISSOURI, DEPT BIOCHEM, 117 SCHWEITZER

HALL/COLUMBIA//MO/65211; UNIV WISCONSIN, DEPT BIOCHEM/MADISON//WI/53706; INDIANA UNIV PURDUE UNIV, DEPT BIOL/INDIANAPOLIS//IN/46202

Journal: PLANT MOLECULAR BIOLOGY, 1993, V21, N6 (MAR), P1011-1021

ISSN: 0167-4412

Language: ENGLISH Document Type: ARTICLE

Abstract: The location of gene expression of the Agrobacterium tumefaciens ipt gene promoter in transgenic tobacco plants was examined using the beta-glucuronidase (GUS) reporter gene. Expression of GUS was detected in every organ and most cell types examined. The highest levels of GUS activity were found in roots. To further examine the transcriptional basis of this broad expression pattern, deletions in the 5' noncoding region of the gene were translationally fused to two promoterless reporter genes, encoding the enzymes chloramphenical acetyl transferase (CAT) and beta-glucuronidase (GUS). Reporter enzyme assays revealed the existence of an upstream segment required for maximal promoter function, the 5' end of which is between -442 and -408 of the P(ipt) ATG codon. This upstream segment is required for maximal levels of GUS expression in roots, but not in other organs, and a tobacco suspension-cultured cell line. The implications of broad ipt expression on the process of crown gall tumorigenesis are discussed.

12/3,AB/98 (Item 29 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
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02342510 Genuine Article#: KV327 Number of References: 51
Title: HORMONAL CONTENT AND SENSITIVITY OF TRANSGENIC TOBACCO AND POTATO
PLANTS EXPRESSING SINGLE ROL GENES OF

AGROBACTERIUM-RHIZOGENES T-DNA (Abstract Available)
Author(s): SCHMULLING T; FLADUNG M; GROSSMANN K; SCHELL J
Corporate Source: UNIV TUBINGEN, LEHRSTUHL ALLGEMEINE GENET, MORGENSTELLE
28/W-7400 TUBINGEN 1//GERMANY/; MAX PLANCK INST ZUCHTUNGSFORSCH/W-5000
COLOGNE 30//GERMANY/; BASF, LANDWIRTSCHAFT VERSUCHSSTN/W-6703
LIMBURGERHOF//GERMANY/

Journal: PLANT JOURNAL, 1993, V3, N3 (MAR), P371-382

ISSN: 0960-7412

Language: ENGLISH Document Type: ARTICLE

Abstract: The expression of single rol genes of the T(L)-DNA of Agrobacterium rhizogenes strain A4 in transgenic tobacco (Nicotiana tabacum L.) and potato (Solanum tuberosum L.) plants alters the internal concentrations of, and the sensitivity to, several plant hormones. The levels of immunoreactive cytokinins, abscisic acid, gibberellins and indole-3-acetic acid were analysed in tissues of the apical shoots, stems, leaves, roots and undifferentiated callus tissue. The addition of the dominant and morphogenetically active rolA, rolB, or rolC genes resulted in alterations in the content of several hormones. rolC overexpression in particular led to an up to fourfold increase in the content of isopentenyladenosine, dihydrozeatin riboside and trans-zeatin riboside-type cytokinins in potato plants. This increase correlated well with different levels of expression of the rolC gene in different transgenic plants. Furthermore it was shown that the dwarfism of P35s-rolC transgenic tobacco and potato plants is correlated with a 28-60% reduction of gibberellic acid Al concentration in apical shoots. Exogenous addition of gibberellic acid completely restored stem elongation in P35S-rolC transgenic plants. Apical shoots of dwarf rolA transgenic tobacco plants also contained 22% less gibberellic acid A1 than control plants, but growth cannot be restored completely by exogenously added gibberellic acid. Similarly, the sensitivity of transgenic tobacco seedlings or callus tissues towards different phytohormone concentrations can be altered by the expression of single rol genes. The overexpression of the rolC gene in seedlings led to an altered response to auxins, cytokinins, abscisic acid, gibberellic acid and the ethylene precursor 1-aminocyclopropane-carboxylic acid. The overexpression of the rolB gene in tobacco calli led to necrosis at lower auxin concentrations than in the wild-type, while other parameters of auxin action, like the induction of cell growth, remained unchanged.

12/3,AB/99 (Item 30 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
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Author(s): STILLER J; NASINEC V; SVOBODA S; NEMCOVA B; MACHACKOVA I
Corporate Source: CZECHOSLOVAK ACAD SCI, INST PLANT MOLEC BIOL, DEPT NITROGEN
FIXAT, BRANISOVSKA 31/CS-37005 CESKE BUDEJOVICE//CZECHOSLOVAKIA/;
CZECHOSLOVAK ACAD SCI, INST EXPTL BOT/CS-16000 PRAGUE 6//CZECHOSLOVAKIA/

Journal: PLANT CELL REPORTS, 1992, V11, N7 (JUL), P363-367

Language: ENGLISH Document Type: ARTICLE

Abstract: Kidney vetch seedlings were induced to form hairy roots by inoculating their mesocotyls with the wild-type strain 15834 of Agrobacterium rhizogenes or with the A. tumefaciens strain C58Cl containing a binary vector system (the pRiA4b as a helper and the vector pCB1346 bearing a pTiC58-derived isopentenyl transferase gene (ipt, cytokinin biosynthetic gene) under control of its native regulatory sequences). Transgenic lines of three distinct phenotypes were selected: (i) Typically, the pRil5834-transformed tissues were stabilized in vitro and maintained for long periods as aseptic, fast-growing, hormone-independent, plagiotropic hairy root cultures which never regenerated shoots and lost the ability to synthesize opines. Their genomic DNA contained both the T(L) - and the T(R) -DNA. (ii) One of the HR-lines transgenic for the T-DNA of pRi15834 (named 52AV34) started to regenerate spontaneously into teratomous shoots. The shoots were found to produce opines and both the T(L) and T(R) parts of T-DNA were found to be partly deleted and/or rearranged. They contained phytohormones in similar levels as those found in seed-born shoots. (iii) A practically identical morphogenic response as in the line 52AV34 was observed in the clone 27AV46. However, its shooty, dark-green, slow-growing teratomas were proven to be kanamycin-resistant, opine-producing, and double-transformed by the pRiA4b sequences and the ipt gene. They over-produced auxins as well as cytokinins (mainly indoleacetylaspartic acid and ribosides of zeatin and isopentenyladenine).

12/3,AB/100 (Item 31 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
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01048832 Genuine Article#: FR384 Number of References: 26
Title: EFFECTS OF THE INTRODUCTION OF AGROBACTERIUM-TUMEFACIENS T-DNA
IPT GENE IN NICOTIANA-TABACUM-L CV PETIT HAVANA SR1 PLANT
-CELLS (Abstract Available)

Author(s): BEINSBERGER SEI; VALCKE RLM; DEBLAERE RY; CLIJSTERS HMM; DEGREEF JA; VANONCKELEN HA

Corporate Source: UNIV INSTELLING ANTWERP, DEPT BIOL/B-2610
WILRIJK/BELGIUM/; LIMBURGS UNIV CENTRUM, DEPT SBM/B-3590
DIEPENBEEK//BELGIUM/; STATE UNIV GHENT, GENET LAB/B-9000 GHENT//BELGIUM/
Journal: PLANT AND CELL PHYSIOLOGY, 1991, V32, N4, P489-496
Language: ENGLISH Document Type: ARTICLE

Abstract: Transformation of tobacco leaf discs with the 'cytokinin' ipt gene yielded several transgenic callus tissue lines, respective to the kind of ipt construction present in the A. tumefaciens cointegrates. Those calli containing an active ipt gene were able to grow hormone-autotrophically and showed an increased endogenous cytokinin level in comparison with controls. Analysis of endogenous IAA level did not allow any quantitative correlation with the cytokinin content. However, a minimal level of auxin seems to be necessary to obtain hormone-autotrophic growth. Exogenously supplied NAA significantly reduced the endogenous cytokinin content without modifying growth characteristics.

The varying chlorophyll content in the different callus lines elicited the study of the ultrastructure of the plastids. The controls contained small plastids, often filled with starch or accumulated vesicles that did not allow observation of the internal membrane system. The 'Pssu-ipt' line, having a higher cytokinin content, showed plastids with an internal membrane system consisting of stroma and grana thylakoids, but this structure was lost during subculture. Swollen thylakoids appeared, the amount of starch was reduced and vesicles were accumulating.

12/3,AB/101 (Item 1 from file: 50) DIALOG(R)File 50:CAB Abstracts (c) 2002 CAB International. All rts. reserv.

CAB Accession Number: 981602461

role of a biosynthetic cytokinin gene in regulating expression of a class of pathogenesis-related protein genes in tobacco plants.

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Institute of Botany, Academia Sinica, Beijing 100093, China.

Acta Botanica Sinica vol. 38 (11): p.870-874

Publication Year: 1996

ISSN: 0577-7496 --

Language: Chinese Summary Language: english

Document Type: Journal article

Expression of basic chitinase, beta -1, 3-glucanase, osmotin and extensin were studied in subcultured seedlings of tobacco cv. Wisconsin 38 growing on MS medium. Total RNA was isolated from tobacco tissues and fractionated on formaldehyde 1.5% agarose gels, blotted onto nylon membranes and hybridized against radioactive probes. Results showed that these 4 genes were regulated in a developmental and organ-specific manner. In transgenic fascicular shoots which contain the active cytokinin

biosynthetic gene (ipt) from Agrobacterium expression of these 4 genes was co-regulated by overproduction of endogenous cytokinins. Heat shock also decreased steady-state levels tumefaciens, of the four mRNAs. 14 ref.

12/3,AB/102 (Item 2 from file: 50) DIALOG(R)File 50:CAB Abstracts (c) 2002 CAB International. All rts. reserv.

03450465 CAB Accession Number: 971610583

Properties of plasma membranes of Phsp 70-ipt transformed tobacco (Nicotiana tabacum).

Bultynck, L.; Geuns, J. M. C.; Ginkel, G. van; Caubergs, R. J.

Ruca, Groenenborgerlaan 171, 2020 Antwerp, Belgium.

Phytochemistry vol. 45 (7): p.1337-1341

Publication Year: 1997

ISSN: 0031-9422 --Language: English

Document Type: Journal article

Application of 10 successive daily heat shocks reduced the growth of control tobacco (Nicotiana tabacum cv. Petit Havana SR1) plants by about 15%; for Phsp 70-ipt transformed plants this was about 48%. The shoot diameter of these ipt-transformed plants increased by about 75%. In addition, in heat shock treated iptplants (IPT -HS) the upper lateral buds grew out due to a reduction of apical dominance. The older leaves of IPT-HS plants had a higher chlorophyll content. In spite of the observed effects due to a higher endogenous cytokinin content in the IPT-HS plants, no significant changes were observed on the plasma membrane fatty acid composition, nor on its fluidity as determined from the steady-state fluorescence anisotropy of DPH. Only a minor change in the plasma membrane free sterol composition was found as evidenced by a 20% decrease in the stigmasterol to sitosterol ratio in IPT-HS, indicative for a possible anti-senescence effect of enhanced endogenous cytokinins , but without significant effects on the plasma membrane function. 26 ref.

12/3,AB/103 (Item 3 from file: 50) DIALOG(R)File 50:CAB Abstracts

(c) 2002 CAB International. All rts. reserv.

03400637 CAB Accession Number: 970607533

Auxin-cytokinin interactions in transgenic expressing the A. tumefaciens ipt, iaaaM and iaaaH genes. Eklof, S.

Department of Forest Genetics and Plant Physiology, Swedish University of Agricultural Sciences, S-901 83 Umea, Sweden.

Acta Universitatis Agriculturae Sueciae - Silvestria

(No. 15): 45 pp.

Publication Year: 1996

Publisher: Swedish University of Agricultural Sciences -- Uppsala, Sweden

ISBN: 91-576-5219-8

Language: English Summary Language: swedish

Document Type: Thesis

The thesis is based on six papers (included as an appendix), and presents results from studies on how plant morphology is influenced by cytokinins and auxins, their metabolism and interactions. The studies were mainly conducted with tobacco plants (Nicotiana tabacum cultivars), with hybrid aspen (Populus tremula x P. tremuloides) chosen transformation with the promoter-less ipt gene. Protein synthesis in the cambial region of Scots pine (Pinus sylvestris) shoots during reactivation was also studied; understanding and control of growth and development of commercial Swedish forest species such as Scots pine is a long-term aim. 8 pp. of ref.

12/3,AB/104 (Item 4 from file: 50) DIALOG(R) File 50:CAB Abstracts (c) 2002 CAB International. All rts. reserv.

03392367 CAB Accession Number: 971606920

The expression of GUS gene driven by T-cyt promoter in transgenic tobacco and potato.

Ma Mi; Zhou DaFeng; Guo Yang; Kuang TingYun; Tang PeiSong; Lin ZhongPing Institute of Botany, Academia Sinica, Beijing 100093, China.

Acta Botanica Sinica vol. 38 (3): p.169-173

Publication Year: 1996

ISSN: 0577-7496 --

Language: Chinese Summary Language: english

Document Type: Journal article

The location of GUS (widA) gene expression under control of the T-cyt gene promoter (gene 4 of T-DNA encoding isopentenyl transferase) (from Agrobacterium tumefaciens) was examined by biochemical assays in transgenic tobacco (Nicotiana tabacum cv. W38) and potato (Solanum tuberosum cv. Desiree) plants. Results showed that T-cyt was expressed in roots, stems, leaves and buds, and the highest levels of GUS activity were found in tobacco stems during axillary bud initiation and in potato buds on tubers. Levels of expression

were also high in wounded leaves of transgenic potato. GUS expression in transgenic tobacco stems was induced cytokinin treatment but not by auxin treatment, indicating that the T-cyt promoter might be selectively induced by exogenous plant

12/3,AB/105 (Item 5 from file: 50) DIALOG(R)File 50:CAB Abstracts (c) 2002 CAB International. All rts. reserv.

03379177 CAB Accession Number: 971605619 Growth pattern, tuber formation and hormonal balance in in vitro potato plants carrying ipt gene.

Machackova, I.; Sergeeva, L.; Ondrej, M.; Zaltsman, O.; Konstantinova, T.; Eder, J.; Ovesna, J.; Golyanovskaya, S.; Rakitin, Y.; Aksenova, N.

De Montfort University, Norman Borlaug Institute for Plant Sciences, Institute of Experimental Botany, Academy of Sciences of the Czech Republic, Ke dvoru 15, 166 00 Prague 6, Czech Republic.

Plant Growth Regulation vol. 21 (1): p.27-36

Publication Year: 1997

ISSN: 0167-6903 --Language: English

Document Type: Journal article

Nodal cuttings of in vitro grown potato (Solanum tuberosum cv. Miranda) plants were transformed by a vector plasmid carrying ipt (isopentenyl transferase) gene of Agrobacterium tumefaciens. From the initial teratoma stage, 5 clones of transgenic plants were obtained, which displayed, in varying degree, shortening of internodes, decreased leaf size, decreased apical dominance and poor rooting. In addition, two of the clones showed increased stolon and tuber formation. In all these clones the endogenous level of free cytokinins (CKs) was increased by 40% to almost 300%. Also, free IAA level was increased, but to a lower degree; the highest increase was 160%. Applied kinetin or (1 mg l-1) strongly suppressed root and tuber formation in two of the clones, although they did not affect or even stimulated these processes in control plants. For control plants the minimal medium sucrose concentration necessary for tuber initiation was 6%, whereas in one clone 2% was sufficient. Differences in the distribution of endogenous CKs and IAA was observed between one clone and control plants. CK content was highest in transgenic plants in stems and in controls in leaves. In one clone, the abscisic acid level was significantly increased in comparison to the control throughout the cultivation period. Ethylene formation was strongly increased during the first week after subcultivation, and later on the difference between transgenic and control plants rapidly diminished. Reactions of plants of one clone to red (RL) and blue light (BL) were similar to reactions of control plants: in RL plants were tall and thin with stunted leaves, in BL they had a teratoma-like appearance and formed a very high number of tubers. The role of hormones in these changes in growth and tuber formation is discussed. 36 ref.

12/3,AB/106 (Item 6 from file: 50) DIALOG(R) File 50:CAB Abstracts (c) 2002 CAB International. All rts. reserv.

03305917 CAB Accession Number: 961611504

Effect of cytokinin on alkaloid accumulation in periwinkle callus cultures transformed with a light-inducible ipt gene.

Garnier, F.; Carpin, S.; Label, P.; Creche, J.; Rideau, M.; Hamdi, S. EA 1370, Laboratoire de Biologie Cellulaire et Biochimie Vegetale, Faculte de Pharmacie, 31 avenue Monge, 37200 Tours, France.

Plant Science (Limerick) vol. 120 (1): p.47-55

Publication Year: 1996

ISSN: 0168-9452 --Language: English

Document Type: Journal article

The effect of cytokinins on accumulation of indole alkaloids in periwinkle (Catharanthus roseus) callus cultures was investigated. Firstly, it was found that exogenously-applied cytokinin increased the ajmalicine and serpentine content of untransformed callus culture obtained from cotyledons. Secondly, periwinkle cotyledons were transformed with the isopentenyl transferase (ipt) gene under the control of a light-inducible promoter and two transformed callus lines used in order to investigate whether endogenously-produced cytokinin could also increase the alkaloid production. It was found that the ipt-transgenic tissues accumulated higher levels of

isopentenyl transferase transcripts as well as zeatin riboside, even under non-inductive condition, but lower concentration of alkaloids compared to that of untransformed tissues. A 28 kDa polypeptide whose accumulation was previously found to be associated with alkaloid production in a periwinkle cell suspension was also present in the non-transformed tissue and its level was increased in parallel to the cytokinin -enhanced alkaloid production. Neither light induction condition, nor exogenous cytokinin treatment led to the increase of the 28 kDa polypeptide accumulation in the transformed tissues. All these data show that endogenously-produced cytokinin does not mimic the effect of exogenously-applied cytokinin on the alkaloid production in periwinkle calli. 34 ref.

12/3,AB/107 (Item 7 from file: 50)
DIALOG(R)File 50:CAB Abstracts
(c) 2002 CAB International. All rts. reserv.

03305846 CAB Accession Number: 961611433

Transgenic periwinkle tissues overproducing cytokinins do not accumulate enhanced levels of indole alkaloids.

Garnier, F.; Label, P.; Hallard, D.; Chenieux, J. C.; Rideau, M.; Hamdi, S.

EA 1370, Laboratoire de Biologie Cellulaire et Biochimie Vegetale, Faculte de Pharmacie, 31 avenue Monge, 37200 Tours, France.

Plant Cell, Tissue and Organ Culture vol. 45 (3): p.223-230 Publication Year: 1996

ISSN: 0167-6857 --Language: English

Document Type: Journal article

Cytokinins play a critical role in several aspects of plant growth, metabolism and development. It has been reported previously that adding cytokinins to the culture medium of a suspension-cultured cell line of periwinkle (Catharanthus roseus) increased the accumulation of indole alkaloids. Studies were conducted to investigate the effects of exogenously applied cytokinins and elevated levels of endogenous cytokinins on indole alkaloid production. An Agrobacterium tumefaciens strain yielding a plasmid with the isopentenyl transferase gene under control of its own promoter was used. Co-culture of suspension cells with the bacteria caused a severe stress response leading to cell necrosis; thus, this material was not transformed. However, periwinkle cotyledons were successfully transformed. It was confirmed that callus cultures generated from the isopentenyl

transferase-transgenic cotyledons accumulated high cytokinin concentrations. Treating normal callus cultures (generated from untransformed cotyledons) with cytokinins enhanced their alkaloid production. In contrast, the enhanced concentration of endogenous cytokinins in transgenic calluses did not increase indole alkaloid production, and thus did not mimic the effect of exogenously applied cytokinins. Hypotheses to explain this discrepancy are discussed.

12/3,AB/108 (Item 8 from file: 50)
DIALOG(R)File 50:CAB Abstracts
(c) 2002 CAB International. All rts. reserv.

03305776 CAB Accession Number: 961611363

Effect of alien ipt gene on hormonal concentrations of plants.

Makarova, R. V.; Borisova, T. A.; Machackova, I.; Kefeli, V. I. Timiryazev Institute of Plant Physiology, Russian Academy of Sciences, ul. Botanicheskaya 35, Moscow 127276, Russia.

Plant hormone signal perception and transduction: Proceedings of

the International Symposium, Moscow, Russia, September 4-10, 1994.
Conference Title: Plant hormone signal perception and transduction:
Proceedings of the International Symposium, Moscow, Russia, September 4-10, 1994.
p.171-173
Publication Year: 1996
Editors: Smith, A. R.; Berry, A. W.; Harpham, N. V. J.; Moshkov, I. E.;
Novikova, G. V.; Kulaeva, O. N.; Hall, M. A.

Publisher: Kluwer Academic Publishers -- Dordrecht, Netherlands ISBN: 0-7923-3768-9
Language: English

Document Type: Conference paper

Transgenic plants carrying the isopentenyl transferase gene (ipt) and normal tobacco plants (Nicotiana tabacum) were analysed to compare their phytohormone status. Total cytokinin (zeatin, zeatin riboside, isopentenyladenine and isopentenyladenosine) level and free IAA content were always higher in shoots regenerated from transgenic cultures although the concentrations were lower in roots. In transgenic plants, IAA-oxidase activity was lower and the concentration of its protectant chlorogenic acid was increased. Transgenic plants also contained lower concentrations of abscisic acid. 14 ref.

12/3,AB/109 (Item 9 from file: 50)
DIALOG(R)File 50:CAB Abstracts
(c) 2002 CAB International. All rts. reserv.

Photosynthesis in transgenic Pssu-ipt tobacco plants as affected by water stress.

Synkova, H.; Pospisilova, J.; Valcke, R.

Institute of Experimental Botany, Na Karlovce 1a, 160 00 Prague 6, Czech Republic.

Photosynthesis: from light to biosphere. Volume IV. Proceedings of the Xth International Photosynthesis Congress, Montpellier, France, 20-25 August 1995.

Conference Title: Photosynthesis: from light to biosphere. Volume IV. Proceedings of the Xth International Photosynthesis Congress, Montpellier, France, 20-25 August 1995.

p.561-564

Publication Year: 1995 Editors: Mathis, P.

Publisher: Kluwer Academic Publishers -- Dordrecht, Netherlands

ISBN: 0-7923-3860-X Language: English

Document Type: Conference paper

In order to determine if water deficit which occurs in ipt (
isopentenyl transferase (which catalyses the initial step in
cytokinin biosynthesis)) transgenic plants especially in the
light originates from suppression of the root system or from effects on
stomatal opening, studies were made of plants of Nicotiana tabacum
cv. Petit Havana SR1 transformed for the ipt gene under the control
of the Pssu promoter. Compared to wild type plants, transgenic
plants exhibited: (1) an approximately 6-fold increase in endogenous
cytokinin content but the same or lower abscisic acid content; (2)
severely affected electron transport around photosystem (PS) I but not
PSII; (3) low stomatal conductance and water potential mainly in mature
and older leaves, probably the result of a poor root system; (4) lower
photosynthesis and higher photorespiration due possibly to closed stomata
and permanent water and CO2 deficits; and (5) no marked disturbances in
PSII functioning, indicating the presence of very efficient water stress
defence mechanisms. 7 ref.

12/3,AB/110 (Item 10 from file: 50) DIALOG(R) File 50:CAB Abstracts (c) 2002 CAB International. All rts. reserv.

03242698 CAB Accession Number: 961606308

Agrobacterium-mediated transformation of commercial mints.

Berry, C.; Eck, J. M. van; Kitto, S. L.; Smigocki, A.

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Plant Cell, Tissue and Organ Culture vol. 44 (2): p.177-181

Publication Year: 1996

ISSN: 0167-6857 Language: English

Document Type: Journal article

Commercial peppermint (P; Mentha x piperita cv. Black Mitcham), native spearmint (NS; M. spicata) and Scotch spearmint (SS; M. x gracillis (M. xgracilis) cv. Baker) petioles, and orange mint (OM; M. (piperita var.) citrata) leaf discs were cocultivated with a number of Agrobacterium tumefaciens strains. P, SS and OM initiated tumour-like callus tissue on growth regulator-free MS medium after cocultivation with strain A281, a hypervirulent agropine strain containing Ti plasmid pTiBo542. Callus did not initiate from explants cocultivated with strain C58, a virulent nopaline strain, with A136, a plasmidless strain, or from uninoculated controls. A281-derived callus was maintained on growth regulator-free medium in the absence of antibiotics for up to two years with no bacterial outgrowth. No shoots regenerated from any of the tumours on regeneration medium. Five of seven OM callus lines assayed gave a positive signal for agropine. DNA extracted from OM tumour tissue hybridized to a DNA probe specific to the T-DNA region of pTi plasmid. Genomic Southern analysis of DNA from tumours of P and SS indicated that one to a few copies of the T-DNA integrated into the mint chromosomes. PCR amplification of genomic DNA with primers specific for one of the T-DNA encoded genes yielded fragments that, when analysed by restriction enzyme mapping and on Southern blots, corresponded to the cytokinin biosynthesis gene ipt (isopentenyl transferase). These results demonstrate transformation of three species of mint and the potential for using A. tumefaciens to transfer economically important genes into commercial mint cultivars. 11 ref.

12/3,AB/111 (Item 11 from file: 50) DIALOG(R)File 50:CAB Abstracts (c) 2002 CAB International. All rts. reserv.

03191981 CAB Accession Number: 961602281

The pattern of cytokinin content in transgenic and wild-type tobacco seedlings as affected by heat shock.

Veselov, S. Y.; Kudoyarova, G. R.; Mustafina, A. R.; Valcke, R.

Institute of Biology, Bashkir Scientific Center, Russian Academy of Sciences, pr. Oktyabrya 69, Ufa, Bashkortostan 450054, Russia. Russian Journal of Plant Physiology vol. 42 (5): p.617-620

Publication Year: 1995

ISSN: 1021-4437 Language: English

Document Type: Journal article

The pattern of the endogenous cytokinin content was monitored during the day in the shoots of transgenic tobacco (Nicotiana tabacum)

plants containing a heat-inducible ipt gene responsible for transferase synthesis. Heating plants at 40 deg C for 1 h yielded an increase in endogenous transgenic cytokinins, as compared to the normal level in the plants kept at 24 deg C for the whole period. However, this increase was not permanent, as after 5 h following heat-shock treatment, there was essentially no difference in **cytokinin** content between heated and untreated **plants**. In the shoots of wild-type tobacco, heat shock activated the processes diminishing **cytokinin** concentration, which are the typical **plant** response to heat shock. When such a response also manifests itself in transgenic **plants**, it can cause a transient **cytokinin** accumulation after heat shock treatment. 12 ref.

12/3,AB/112 (Item 12 from file: 50)
DIALOG(R)File 50:CAB Abstracts
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03164555 CAB Accession Number: 961600117

Agrobacterium-mediated transformation of the apple cultivar Granny Smith.

Trifonova, A.; Savova, D.; Ivanova, K.

Institute of Genetic Engineering, 2232 Kostinbrod, Bulgaria.

Progress in temperate fruit breeding. Proceedings of the Eucarpia Fruit Breeding Section Meeting, Wadenswil/Einsiedeln, Switzerland, 30 August to 3 September, 1993.

Conference Title: Progress in temperate fruit breeding. Proceedings of the Eucarpia Fruit Breeding Section Meeting, Wadenswil/Einsiedeln, Switzerland, 30 August to 3 September, 1993.

p.343-347

Publication Year: 1994

Editors: Schmidt, H.; Kellerhals, M.

Publisher: Kluwer Academic Publishers -- Dordrecht, Netherlands

ISBN: 0-7923-2947-3 Language: English

Document Type: Conference paper

An efficient adventitious shoot regeneration system from leaf segments the apple cultivar Granny Smith was developed. Regenerants in of sufficient frequency were obtained under the optimal conditions in presence of 3 mg BA, 2 mg 2iP and 0.2 mg NAA/litre. Putative transgenic plants were regenerated from leaf segments that were co-cultivated with disarmed C58 Agrobacterium tumefaciens strain containing either of the following binary plasmids: pGV2449 or pGV2492. The chimaeric marker gene for neomycin phosphotransferase II (nptII) and ipt genes (encoding for isopentenyl transferase, the first enzyme in the cytokinin biosynthetic pathway) were integrated in both plasmid derivatives. Seven putative transgenic plants were obtained on the selective medium containing 50 micro g/ml kanamycin after transformation with pGV2449. The **expression** and integration of nptII marker gene was detected in leaves of the **plants**. Rooting of the propagated plants was only achieved in presence of anticytokinin substance, 4-substituted-triazolo (4,5,d) pyrimidine and 0.5 mg IBA/litre. 11 ref.

12/3,AB/113 (Item 13 from file: 50)
DIALOG(R)File 50:CAB Abstracts
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03060158 CAB Accession Number: 951608688

Cytokinin involvement in the control of coumarin accumulation in Nicotiana tabacum. Investigations with normal and transformed tissues carrying the isopentenyl transferase gene.

Hamdi, S.; Creche, J.; Garnier, F.; Mars, M.; Decendit, A.; Gaspar, T.; Rideau, M.

Laboratoire de Biologie Cellulaire et Biochime Vegetale, Faculte de Pharmacie, EA 1370, 37200 Tours, France.

Plant Physiology and Biochemistry (Paris) vol. 33 (3): p.283-288

Publication Year: 1995

ISSN: 0981-9428 --Language: English Document Type: Journal article

The effects of cytokinins on accumulation of the coumarin scopolin in tobacco tissues were investigated. Leaf discs were transformed with the Agrobacterium tumefaciens ipt gene under control of either its native promoter or a light-inducible (Rubisco (ribulose-bisphosphate carboxylase/oxygenase)) promoter. Several shoot cultures were isolated, from which ipt transgenic callus cultures were initiated. Leaves from all the ipt transgenic shoot cultures (grown in light) accumulated a high level of scopolin, whereas control (untransformed) leaves did not. Callus cultures carrying ipt under the control of its own promoter accumulated higher contents of scopolin as compared with calluses, irrespective of light or dark conditions. untransformed Dark-grown callus cultures carrying ipt under the control of the light-inducible promoter accumulated scopolin to levels comparable with untransformed calluses. Transferring transgenic calluses from dark to light, or adding a cytokinin to the culture medium resulted in an increase of the scopolin content. Exogenously applied cytokinin also increased the scopolin content of untransformed callus cultures. These data indicated that cytokinins control coumarin accumulation, and that enhanced levels of endogenous cytokinins could mimic the effect of exogenous cytokinins on coumarin pathway in tobacco tissues. 27 ref.

12/3,AB/114 (Item 14 from file: 50)
DIALOG(R)File 50:CAB Abstracts
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02924669 CAB Accession Number: 941610347

Cytokinin accumulation and action: biochemical, genetic, and molecular approaches.

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Plant Science Institute, Department of Biology, University of Pennsylvania, Philadelphia, PA 19104-6018, USA.

Annual Review of Plant Physiology and Plant Molecular Biology vol. 45 p.173-196

Publication Year: 1994

ISSN: 1040-2519 --Language: English

Document Type: Journal article

Progress in identifying the genes and gene products involved in cytokinin control of growth and development is reviewed. Biochemical approaches are considered under the headings cytokinin biosynthesis, and **cytokinin** receptors. metabolism Genetic cytokinin study of cytokinins have made use of approaches to the cytokinin accumulation mutants and cytokinin response mutants. The molecular approaches discussed include those used in the study of and transgenic cytokinin control of gene expression ipt (encoding expressing the gene isopentenyltransferase) from Agrobacterium tumefaciens. 134 ref.

12/3,AB/115 (Item 15 from file: 50)
DIALOG(R)File 50:CAB Abstracts
(c) 2002 CAB International. All rts. reserv.

02899052 CAB Accession Number: 941609077

Genetic transformation of some poplar clones.

Original Title: Transformarea gentica a unor clone de plop. Ionita, L.

Institutul de Cercetari si Amenajari Silvice, 72902 Bucharest, Romania.

Probleme de Genetica Teoretica si Aplicata vol. 25 (2): p.99-111

Publication Year: 1993 -

Language: Romanian Summary Language: english

Document Type: Journal article

The clones used were 717 1B4 (Populus tremula x P. alba (P. canescens)), Beaupre and Boelare (both P. trichocarpa x P. deltoides (P. interamericana)) and Ogy (P. deltoides x P. nigra (P. canadensis)). The genetic transformation was by coculture with Agrobacterium tumefaciens. Different vectors were tested. Clone 717 1B4 was successfully transformed using plasmid p35SASOM3C carrying the gene for O-methyltransferase (involved in lignin synthesis). The other 3 clones showed no regeneration when transformed with the same constructs. A cloning strategy was developed for the Tmr (Ipt) gene with the PRI-a promoter. This gene codes for an enzyme involved in the synthesis of cytokinins and when introduced into the plant genome conditions an acceleration of growth. Earlier tests involving transformation with this gene led to an abnormal development of transformed plants; hence the use in this case of an inducible promoter (PRI-a), which allows control of gene expression in the plant. 8 ref.

12/3,AB/116 (Item 16 from file: 50)
DIALOG(R)File 50:CAB Abstracts
(c) 2002 CAB International. All rts. reserv.

Morphometric analysis of the growth of Phsp 70-ipt transgenic tobacco plants.

Loven, K. van; Beinsberger, S. E. I.; Valcke, R. L. M.; Onckelen, H. A. van; Clijsters, H. M. M.

Limburgs Universitair Centrum (LUC), Department SBG, Universitaire Campus, 3590 Diepenbeek, Belgium.

Journal of Experimental Botany vol. 44 (268): p.1671-1678

Publication Year: 1993

ISSN: 0022-0957 --Language: English

Document Type: Journal article

The effect of introducing a supplementary <code>ipt</code>-gene into the genome of Nicotiana tabacum cv. Petit Havana SR1 was studied. The <code>ipt</code>-gene, accounting for the biosynthesis of <code>cytokinins</code>, was coupled to the heat-inducible hsp70 promoter from Drosophila melanogaster. The influence of the hormonal changes involved was examined as well as the effects of the in vitro growth conditions used for selecting transformed <code>plants</code>

and the heat treatment to induce <code>ipt-gene expression</code>. The phenotype of the <code>plants</code> was determined by the tissue sensitivity to three factors: (1) heat treatment reduces stem elongation and diameter growth; (2) in vitro pre-cultivation also reduces stem elongation; and (3) <code>expression</code> of the <code>ipt-gene</code> stimulates diameter growth, induces debudding in the axillary shoots and <code>inhibits</code> root development. In addition, axillary bud development indicates that in vitro cultivation, implying a stress condition, affects hsp70-<code>ipt</code> gene <code>expression</code>. 26 ref.

12/3,AB/117 (Item 17 from file: 50)
DIALOG(R)File 50:CAB Abstracts
(c) 2002 CAB International. All rts. reserv.

02898514 CAB Accession Number: 941608524

Attempts to elucidate the molecular mechanism of genetic tumors in Nicotiana.

Feng, X. H.; Kung, S. D.

Center for Agricultural Biotechnology and Department of Botany, University of Maryland, College Park, MD 20742, USA.

Institute of Botany, Academia Sinica Monograph Series (No. 13): p.35-46 Publication Year: 1993

ISSN: 0258-5170 --

Language: English

Document Type: Journal article

The tumorous amphidiploid hybrid (GGLL wild type) of N. glauca x N. langsdorffii, a non-tumorous mutant (GGLL mutant), and the parental species were used to study the molecular and physiological mechanisms underlying spontaneous genetic tumorigenesis. Endogenous levels of cytokinins in various tissues of all 4 genotypes were measured in immunoassays. Tumours contained relatively higher level of cytokinin than other tissues. The non-tumorous mutant exhibited a shooty morphology, indistinguishable from that of wild type genetic tumours, when it was treated by exogenously-applied cytokinins or transformed with an Agrobacterium tumefaciens Ti T-DNA gene (ipt) encoding isopentenyltransferase, an enzyme involved in the biosynthesis of cytokinin. This altered phenotype of the transformed mutant was isopentenyltransferase, an caused by an elevation in the level of cytokinin resulting from the constitutive expression of the ipt gene. The spatial and temporal regulation of the Ng rol (N. glauca genomic genes homologous to the A. rhizogenes Ri rol genes) gene expression was also examined in genetic tumors. The expression of Ng rolC was higher in tumours than in normal tissues, suggesting that Ng rolC, which may have a similar function as Ri rolC to release free cytokinins from their conjugated forms, might play an important role in genetic tumour formation and/or maintenance. In conclusion, it seems that genetic tumours were caused, at least in part, by elevated levels of free cytokinin in interspecific hybrids. Furthermore, to identify other regulators of tumour induction and growth, PCR (polymerase chain reaction) was used to isolate protein kinase sequences from Nicotiana. RNA blot analyses showed that transcripts of 4 isolated kinase genes accumulated differentially during genetic tumour induction. Transcription of one protein kinase, named NIPK2, increased during tumour induction, while other kinase transcripts showed little change during the induction period. Thus, protein kinases may play a very critical regulatory role in **plant** hormone-mediated genetic tumorigenesis in Nicotiana. 64 ref.

12/3,AB/118 (Item 18 from file: 50) DIALOG(R)File 50:CAB Abstracts (c) 2002 CAB International. All rts. reserv.

02776801 CAB Accession Number: 931644425

Morphological characteristics and phytohormone content of $\ensuremath{\text{\textbf{ipt}}}$ -transgenic tobacco.

Beinsberger, S. E.; Clijsters, H. M.; Valcke, R. L.; Onckelen, H. van Department of SBM, Limburgs Universitait Centrum, 3590 Diepenbeek, Belgium.

Conference Title: Progress in plant growth regulation. Proceedings of the 14th International Conference on Plant Growth Substances, Amsterdam, Netherlands, 21-26 July 1991

p.738-745

Publication Year: 1992

Editors: Karssen, C. M.; Loon, L. C. van; Vreugdenhil, D.

Publisher: Kluwer Academic Publishers -- Dordrecht, Netherlands

ISBN: 0-7923-1617-7 Language: English

Document Type: Book chapter

Phytohormone content and morphological characteristics were analysed in plant material derived from Nicotiana tabacum cv. Petit Havana SR1 leaf discs transformed with the Agrobacterium tumefaciens T-DNA ipt

gene using recombinant Ti-plasmids pGV2492 and pGV2488. Data on the cytokinin content and cytokinin: auxin ratio are provided for (1) transgenic calluses; (2) transgenic regenerants; (3) transgenic grafts (with transgenic shoots sandwiched in a vertical incision of a decapitated untransformed tobacco plant) and (4) transgenic seedlings. 7 ref.

12/3,AB/119 (Item 19 from file: 50)
DIALOG(R)File 50:CAB Abstracts
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Transgenic **plants** and transgenic **plant** mosaics for the **expression** of pathogen derived genes able to affect phytohormone activity.

Spena, A.; Estruch, J. J.; Aalen, R. D.; Prinsen, E.; Parets-Soler, A.; Nacken, W.; Sommer, H.; Chriqui, D.; Grossmann, K.; Onckelen, H. van; Schell, J.

Max-Planck-Institut fur Zuchtungsforschung, 5000 Koln 30, Germany. Conference Title: Progress in plant growth regulation. Proceedings of the 14th International Conference on Plant Growth Substances, Amsterdam, Netherlands, 21-26 July 1991

p.724-730

Publication Year: 1992

Editors: Karssen, C. M.; Loon, L. C. van; Vreugdenhil, D.

Publisher: Kluwer Academic Publishers -- Dordrecht, Netherlands

ISBN: 0-7923-1617-7 Language: English

Document Type: Book chapter

Information on phytohormone activity in genetically engineered plants containing pathogen derived genes is collated on the basis of studies on (1) genetic mosaics for cytokinin synthesis (ipt

gene from Agrobacterium tumefaciens), (2) genetic mosaic for the expression of the rolC gene of A. rhizogenes, (3) plants transgenic for the IAA lysine synthetase (iaaL) gene of Pseudomonas savastanoi, and (4) plants transgenic for tapetum specific expression of the rolB gene. Most studies were made with tobacco. 14 ref.

12/3,AB/120 (Item 20 from file: 50)
DIALOG(R)File 50:CAB Abstracts
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02762167 CAB Accession Number: 931642872

Modulation of chloroplast gene expression in transgenic plants of tobacco following changes in the phytohormone balance.

Yusibov, V. M.; Pak Chun Ir; Andrianov, V. M.; Piruzyan, E. S. Conference Title: 1 Vsesoyuznyi simpozium "Novye metody biotekhnologii rastenii", Pushchino, 20-22 noyabrya, 1991: Tezisy dokladov.

p.50-51, 152-153

Publication Year: 1991

Publisher: -- Pushchino, Russia

Language: English; Russian

Document Type: Miscellaneous

Transgenic plants of 2 types were produced: containing the Escherichia coli glucose-6-phosphate isomerase gene xyl and the cytokinin synthesis gene ipt from Agrobacterium tumefaciens Ti-plasmid T-DNA. Analysis of plants of both types showed an increase in the content of cytokinins. Northern blot hybridization, which was used to assess accumulation of mRNA of the rbcL gene coding for the large subunit of ribulose-bisphosphate carboxylase, showed an increased content of this mRNA both in the transgenic plants and after treatment with exogenous cytokinin. Changes in the content of mRNAs of some other chloroplast genes in the transgenic plants were studied, e.g. psbA, proB, several ndh genes and the gene for 23S rRNA.

12/3,AB/121 (Item 21 from file: 50)
DIALOG(R)File 50:CAB Abstracts

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02606870 CAB Accession Number: 921632055

Progress in cytokinin research.

Kaminek, M.

Institute of Experimental Botany, Czechoslovak Academy of Sciences, 16630 Prague 6, Czechoslovakia.

Trends in Biotechnology vol. 10 (5): p.159-164

Publication Year: 1992

ISSN: 0167-7799 - Language: English

Document Type: Journal article

The subject is reviewed under the headings; biological effects of cytokinins, how cytokinins originate, tRNA as a source of free cytokinins, metabolism of cytokinins (cytokinin

conjugates and cytokinin oxidase), production of cytokinins in transgenic plants, reducing the cytokinin content of plant cells, habituation, how cytokinins act in plant

cells, production of secondary metabolites in transformed plants, and outlook for the future. The potential is highlighted for using cytokinin genes in transgenic plants to increase yield by expression post anthesis. The Agrobacterium tumefaciens ipt

gene has been expressed in transgenic tobacco and Arabidopsis plants in response to stimulation of a heat shock promoter. Antisense ipt genes might be used to reduce levels of cytokinin relative to auxin, thus stimulating rooting and reduction of branching in some ornamental and forest trees. Key future targets are cloning the genes regulating cytokinin mobilization, degradation and inactivation, and cytokinin binding sites. 42 ref.

12/3,AB/122 (Item 22 from file: 50)
DIALOG(R)File 50:CAB Abstracts
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02518502 CAB Accession Number: 921627354

Delayed leaf senescence in tobacco plants transformed with tmr, a gene for cytokinin production in Agrobacterium.

Smart, C. M.; Scofield, S. R.; Bevan, M. W.; Dyer, T. A.

IPSR Cambridge Laboratory, John Innes Centre, Colney, Norwich NR4 7UH,

Annual report, AFRC Institute of **Plant** Science Research, John Innes Institute and Sainsbury Laboratory, 1990.

p.30-32

Publication Year: 1991?

Publisher: IPSR & John Innes Institute -- Norwich, UK

Language: English

Document Type: Annual report

Cytokinin levels in plants can be controlled by activity of the enzyme isopentenyl transferase, which in A. tumefaciens is controlled by gene tmr. The bacterial promoter for tmr was replaced with a soyabean promoter which is activated by heating to 42 deg C thus enabling direct comparison of adjacent tissue. Following heat shock, transformed leaves remained green while surrounding untransformed tissue died. The level of cytokinins increased markedly after a 2 h heat shock and zeatin concentration was 15-20-fold higher in heat shocked tissue 4 h after heat shock. The amount of tmr mRNA detectable by Northern blot analysis remained up to 8 h after heat shock, but was considerably lower by 24 h after heat shock. 6 ref.

12/3,AB/123 (Item 23 from file: 50)
DIALOG(R)File 50:CAB Abstracts
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Full expression of chimeric T-DNA gene 4 constructions in tobacco tissues.

Beinsberger, S. E.; Rudelsheim, P.; Inze, D.; Lijsebettens, M. van; Greef, J. de; Onckelen, H. A. van

Dep. Biology, University of Antwerp, UIA, B-2610 Wilrijk, Belgium. Archives Internationales de Physiologie et de Biochimie vol. 96 (1): p.PP 2

Publication Year: 1988 --

Language: English

Document Type: Conference paper; Journal article

This paper was presented at a meeting of the Belgian Association of Plant Physiology at Liege on 14th Nov. 1987. The Agrobacterium tumefaciens T-DNA gene 4 encodes for isopentenyl-transferase

which catalyses the first step in cytokinin biosynthesis. Chimeric T-DNA gene 4 constructions incorporated in Nicotiana tabacum cv. Petit Havana SR1. in the pGV831 vector were mobilized in A. tumefaciens in the Ti-plasmid vector pGV2260. Since the pGV831 contained a selectable marker (Pnos-nptII) the transformed cells could be selected on kanamycin-containing medium in presence of both cytokinins so that the activity of different chimeric genes in an auxins identical genetic background could be compared. In the control line, which contained only the selectable marker, very low cytokinin amounts were observed. In tobacco calli containing the octopine (pTi C58)-, the nopaline (pTi B6S3) gene 4 coupled to its own non-light inducible promoter, as well as in calli transformed with the chimeric octopine gene 4 coupled to the Pnos promoter, an increase of the endogenous levels of both **cytokinins** and IAA was observed. Consequently all gene 4-containing tissues managed to survive on a phytohormone-deficient medium. Surprisingly low endogenous cytokinin levels were found in light-grown calli containing a chimeric gene 4 construct coupled to the light-inducible Pssu promoter. Growth experiments on phytohormone-deficient media, however, showed that in the light some of the Pssu-gene 4 lines survived whereas in the dark the same lines turned brown and died. 6 ref.

12/3,AB/124 (Item 24 from file: 50)
DIALOG(R)File 50:CAB Abstracts
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02335370 CAB Accession Number: 901617500

Dual promoter of Agrobacterium tumefaciens mannopine synthase genes is regulated by **plant** growth hormones.

Langridge, W. H. R.; Fitzgerald, K. J.; Koncz, C.; Schell, J.; Szalay, A. A.

Plant Molecular Genetics, University of Alberta, Medical Sciences Building, Edmonton, AB T6G 2P5, Canada.

Proceedings of the National Academy of Sciences of the United States of America vol. 86 (9): p.3219-3223

Publication Year: 1989

ISSN: 0027-8424 --Language: English

Document Type: Journal article

Temporal and spatial distribution of mannopine synthase (mas) promoter activity was determined throughout the development of transgenic tobacco plants using bacterial luciferase luxA and luxB as reporter genes. Luciferase activity was determined by luminometry in vitro and visualized by computer-enhanced single-photon video imaging in vivo. The activity of the mas dual promoters increased basipetally in developing plants and was wound-inducible in leaf and stem tissue. Hormone bioassays with isolated plant tissues and tumours deficient in the transferred DNA (T-DNA)-encoded genes iaaM, iaaH and ipt indicated that activity of

the mas dual promoters is regulated by auxin and enhanced by cytokinin in both differentiated and tumorous plant cells. 33 ref.

12/3,AB/125 (Item 25 from file: 50)
DIALOG(R)File 50:CAB Abstracts
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Agrobacterium-mediated transformation of the cultivated strawberry (Fragaria x ananassa Duch.) using disarmed binary vectors.

James, D. J.; Passey, A. J.; Barbara, D. J.

Institute of Horticultural Research, East Malling, Maidstone, Kent, ME19 6BJ, UK.

Plant Science (Limerick) vol. 69 (1): p.79-94

Publication Year: 1990

ISSN: 0168-9452 --Language: English

Document Type: Journal article

Two disarmed Ti-binary vectors of A. tumefaciens were used to produce viable transgenic strawberry plants. Fertile strawberry plants with a normal phenotype were regenerated after transformation with pBIN6, which carries genes for nopaline synthase (nos) and neomycin phosphotransferase (nptII) (conferring kanamycin resistance). The transfer and expression of the 2 genes was confirmed by Southern blot analysis, the detection of nopaline synthase activity in vegetative and reproductive tissues and rooting in vitro in the presence of kanamycin. The nos gene continued to be **expressed** in greenhouse-grown plants many months after removal from in vitro growth conditions. After selfing the RO plants, nos segregated in the R1 progeny according to a 3 : 1 Mendelian ratio. In in vitro germinated seedlings there was complete correlation between the presence of nopaline synthase activity and the ability of leaf segments to produce callus on a medium containing kanamycin. Transgenic clones that exhibited an abnormal phenotype associated with cytokinin overproduction were produced when **plants** were transformed with pSS1, a derivative of pBIN19 carrying both nptII and **ipt** (encoding isopentenyltransferase). Shoots of these clones grew on hormone-free medium, could not be induced to root and their growth was unaffected by the presence of 50 micro g/mlkanamycin in hormone-free media. 36 ref.

12/3,AB/126 (Item 26 from file: 50)
DIALOG(R)File 50:CAB Abstracts
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02271501 CAB Accession Number: 901145882

Agrobacterium tumefaciens 6b genes are strain-specific and affect the activity of auxin as well as cytokinin genes.

Tinland, B.; Huss, B.; Paulus, F.; Bonnard, G.; Otten, L.

Institut de Biologie Moleculaire des Plantes du CNRS, Rue de General Zimmer 12, 67084 Strasbourg, France.

Molecular and General Genetics vol. 219 (1-2): p.217-224

Publication Year: 1989

ISSN: 0026-8925 --Language: English

Document Type: Journal article

The T-region located 6b gene of A. tumefaciens was found to interfere with **cytokinin** effects produced by the contransferred **ipt** gene. The biological activity of 3 different 6b genes were compared: A-6b from Ach5 (octopine biotype I), C-6b from C58 (nopaline, biotype I) and T-6b from Tm4 (octopine, biotype III). Each 6b gene was inserted into a disarmed vector and tested on tobacco stems in co-infection experiments

with the Ach5 cytokinin (ipt) gene (A-ipt). A-ipt /C-6b co-infections produced tumours with shoots, A-ipt /A-6b co-infections green tumours and A-ipt/T-6b co-infections tumours with a necrotic surface. The tumour phenotypes obtained were independent the 6b/A-ipt co-infection ratios, indicating that the str.-specific 6b effects result from the activity of a non-diffusible 6b encoded product. Studies with **ipt**-less Tm4 mutants showed that 6b genes affect other tumour genes besides the ipt gene and pointed to an influence of T-6b on auxin effects resulting from the Tm4 iaa system. T-iaa/T-6b co-infection experiments showed that T-6b did indeed strongly increase tumour formation by the Tm4 iaa genes. The 3 6b genes also have effects which do not require other T-region genes. The complex role of the 6b gene in crown gall induction and Agrobacterium host range is discussed. 37 ref.

12/3,AB/127 (Item 27 from file: 50) DIALOG(R)File 50:CAB Abstracts (c) 2002 CAB International. All rts. reserv.

CAB Accession Number: 901175516

Nucleotide sequence, evolutionary origin and biological role of a rearranged cytokinin gene isolated from a wide host range biotype III Agrobacterium strain.

Bonnard, G.; Tinland, B.; Paulus, F.; Szegedi, E.; Otten, L. Institut de Biologie Moleculaire des Plantes du CNRS, Rue de General Zimmer 12, 67000 Strasbourg, France.

Molecular and General Genetics vol. 216 (2-3): p.428-438

Publication Year: 1989

ISSN: 0026-8925 Language: English

Document Type: Journal article

A DNA fragment with homology to the cytokinin (ipt) gene from biotype I A. tumefaciens str. Ach5 was cloned from the Ti plasmid of the wide host range biotype III A. str. Tm- 4 and sequenced. The fragment contains an intact ipt coding sequence. However, the 3' non-coding region of this ipt gene is rearranged due to a 0.9 kb deletion fusing it to the 3' coding region of the neighbouring gene 6a, most of which was found to be deleted. The Tm-4 ipt gene is strongly related to the partially deleted ipt gene of the limited host range biotype III str Ag162. To test its biological activity, the Tm-4 ipt gene was inserted into a specially constructed, disarmed Ti vector lacking tzs and tested on tobacco, where the rearranged ipt gene induced shoot formation. The cloned Tm-4 ipt gene was mutated with Tn5 and the intact gene on the wild-type Tm-4 Ti plasmid was replaced by the mutated gene. The resulting str. was avirulent on tobacco but normally virulent on the natural host of the wild-type str. Tm-4, grapevine. As the biotype I 6b gene diminishes the effect of a corresponding ipt gene, a larger Tm-4 fragment carrying both the ipt gene and an adjacent 6b-like gene was also tested on tobacco and compared with the Tm-4 ipt fragment alone and with an ipt and 6b/ipt fragment derived from Ach5. The Tm-4 6b gene diminishes the effect of the Tm-4 ipt gene, showing the Tm-4 6b gene to be active as well. The Tm-4 6b/ ipt combination is less effective than the Ach5 combination. These results provide further insight into the molecular basis of the host range differences between limited host range and wide host range biotype III A. strs and show that the WHR cytokinin gene, although active, does not significantly contribute to tumour formation on grapes, the natural host of the WHR biotype III strs. 57 ref.

12/3,AB/128 (Item 28 from file: 50) DIALOG(R)File 50:CAB Abstracts (c) 2002 CAB International. All rts. reserv.

02147694 CAB Accession Number: 891607282

Construction of a heat-inducible chimaeric gene to increase the cytokinin content in transgenic plant tissue.

Schmulling, T.; Beinsberger, S.; Greef, J. de; Schell, J.; Onckelen, H. van; Spena, A.

MPI fur Zuchtungsforschung, 5000 Koln 30, German Federal Republic.

FEBS Letters vol. 249 (2): p.401-406

Publication Year: 1989

ISSN: 0014-5793 Language: English

Document Type: Journal article
The ipt gene of Agrobacterium tumefaciens T-DNA encodes an isopentenyltransferase which causes cytokinin overproduction and developmental alterations in transformed plants. A chimaeric gene, constructed by positioning the **ipt** coding region under the control of the hsp70 gene from Drosophila melanogaster, allowed heat-regulated expression in transgenic plant tissue. Heat-shock treatment of tobacco calluses transgenic for the chimaeric hsipt gene increased the endogenous cytokinin concentration and enabled the calluses to grow on cytokinin-free medium. Transgenic plants regenerated from calluses transformed with the hsipt gene and grown at normal temperatue were phenotypically normal. 21 ref.

12/3,AB/129 (Item 29 from file: 50) DIALOG(R)File 50:CAB Abstracts (c) 2002 CAB International. All rts. reserv.

CAB Accession Number: 891126000

Hormones and the molecular basis of determination in plants. Meins, F, Jr.

Friedrich Miescher-Inst., CH-4002 Basle, Switzerland.

Monograph, British Plant Growth Regulator Group (No. 16): p.19-28 Publication Year: 1987

Language: English

Document Type: Journal article

Studies of variation in the cytokinin requirement of cultured tobacco tissues show that plant cells can undergo potentially reversible, cell-heritable changes in phenotypic expression. The cytokinin -autotrophic state appeared to be stabilized by a positive-feedback loop in which cytokinins or similar cell-division for the cytokinins of similar cell-division factors induced their own biosynthesis. The cytokinin requirement of cultured tobacco cells was regulated at two genetic loci, Hl-1 and Hl-2. The Hl-1 locus could be activated by mutation to have an oncogenic function similar to the isopentenyl transferase locus of the tumour-inducing Ti plasmid. 22 ref.

12/3, AB/130 (Item 30 from file: 50) DIALOG(R)File 50:CAB Abstracts (c) 2002 CAB International. All rts. reserv.

02016733 CAB Accession Number: 881673565

Cytokinin gene fused with a strong promoter enhances shoot organogenesis and zeatin levels in transformed plant cells.

Smigocki, A. C.; Owens, L. D.

Tissue Culture & Molec. Biol. Lab., ARS, USDA, Beltsville, MD 20705,

Proceedings of the National Academy of Sciences of the United States of America vol. 85 (14): p.5131-5135

Publication Year: 1988

ISSN: 0027-8424 --Language: English

Document Type: Journal article isopentenyltransferase (ipt) gene associated cytokinin biosynthesis in plants was cloned from a tumour-inducing plasmid carried by Agrobacterium tumefaciens and placed under the control of promoters of differing activities, the cauliflower mosaic virus 35S promoter and the nopaline synthase promoter. These promoter-gene constructs were introduced into wounded Nicotiana (N. tabacum, N. rustica and N. plumbaginifolia) stems and leaf pieces and cucumber seedlings by A. tumefaciens infection. Shoots were observed at the infection site on all Nicotiana plants (except those of N. rustica) infected with the 35S promoter construct (35S-ipt), whereas only 41% responded similarly to infection with the unmodified gene. Furthermore, shoots were observed 19 days after infection with the 35Sipt and were up to 6 times taller than shoots induced by the native gene. On cucumber, shoots were observed only on galls incited by the 35Sipt construct. These galls were, on average, 7.5 times larger than those incited by the nopaline synthase promoter construct (NOS-ipt) unmodified ipt gene. Zeatin concentrations, were on average, 23 times higher in 35S-ipt zeatinriboside transformed shoots than in ones transformed with the native ipt gene. The results suggested that a more active promoter on the iptgene can enhance or change the morphogenic potential of transformed plant cells by increasing their endogenous cytokinin levels. 41 ref.

12/3,AB/131 (Item 1 from file: 65) DIALOG(R) File 65: Inside Conferences (c) 2002 BLDSC all rts. reserv. All rts. reserv.

02468009 INSIDE CONFERENCE ITEM ID: CN025771495 Expression of IPT Gene in Transgenic Arabidopsis Plants Leads to Ubnormal Accumulation of Cytokinin N-Glucosides Werner, T.; Rupp, H. M.; Schmuelling, T.; Van Onckelen, H. CONFERENCE: Plant physiology-Czech-Slovak conference; 8th (Eighth days of plant physiology) P: 237 Palacky University, 1998 ISBN: 8070678720

LANGUAGE: English DOCUMENT TYPE: Conference Abstracts CONFERENCE LOCATION: Olomouc, Czech Republic CONFERENCE DATE: Jul 1998 (199807) (199807)

12/3,AB/132 (Item 1 from file: 76) DIALOG(R) File 76:Life Sciences Collection (c) 2002 Cambridge Sci Abs. All rts. reserv.

01534198 2631378

Viviparous leaves produced by somatic activation of an inactive cytokinin-synthesizing gene. Estruch, J.J.; Prinsen, F.; Van Onckelen, H.; Schell, J.; Spena, A. MPI Zuechtungsforsch., Carl-von-Linne weg, 10, 5000 Koeln 30, FRG SCIENCE (WASH.). vol. 254, no. 5036, pp. 1364-1367 (1991.) DOCUMENT TYPE: Journal article LANGUAGE: ENGLISH

SUBFILE: Genetics Abstracts

Tobacco plants that are somatic mosaics for expression of a cytokinin-synthesizing gene have viviparous leaves. Such a formation of shoots in an abnormal position represents a significant deviation from the usual organization of the plant body where a central axis produces shoots only in the axis of lateral leaf appendages and according to a precise phyllotactic pattern. This report links vivipary to the expression of a gene whose product is involved in the synthesis of